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                                                      Search time 17695.9 Seconds (without alignments)
16.701 Million cell updates/sec
                                                                                                                                                                                                                                                                       /cgn2_6/ptodata/1/pna/US6006_COMB.seq:
/cgn2_6/ptodata/1/pna/US6007_COMB.seq:
/cgn2_6/ptodata/1/pna/US6008_COMB.seq:
/cgn2_6/ptodata/1/pna/US6009_COMB.seq:
/cgn2_6/ptodata/1/pna/US6011_COMB.seq:
/cgn2_6/ptodata/1/pna/US6011_COMB.seq:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       'ptodata/1/pna/US6001_COMB.seq'
'ptodata/1/pna/US6002_COMB.seq
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ptodata/1/pna/US6005_COMB.seq:
      GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                          /cgn2_6/ptodata/1/pna/PCTUS_COMB.seq:*
/cgn2_6/ptodata/1/pna/US06_COMB.seq:*
                                                                                                                                                   14155048 seqs, 7388405095 residues
                                                                                                                                                                   Total number of hits satisfying chosen parameters:
                                                       October 2, 2001, 21:50:17
                                                                                                                                                                                                                        Maximum Match 100%
Listing first 45 summaries
                                      OM nucleic - nucleic search, using sw model
                                                                                                        1 attectegetgetggtggaa 20
                                                                                                                                                                                                                                                  Pending_Patents_NA_Main:*
                                                                                                                           IDENTITY_NUC
Gapop 10.0 , Gapext 1.0
                                                                                      US-09-757-100B-17
20
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                                                                                                                                                                                                                                                                                                                                                                        Title:
Perfect score:
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/cgn2_6/ptodata/1/pna/US6015_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6015_COMB.seq:*
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/cgn2_6/ptodata/1/pna/US6019_COMB.seq:*
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/cgn2_6/ptodata/1/pna/US6012_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6012_COMB.seq:
/cgn2_6/ptodata/1/pna/US6013_COMB.seq:
/cgn2_6/ptodata/1/pna/US6014_COMB.seq:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

			di			SUMMARIES	
Re	Result No.	Score	Query Match	Length 1	DB	QI.	Description
	, et c	20	100.0		-1-	- F	17, A
	4 (*)	200	100.0	200	20	-09-757-100	Segmence 17, Appl
υ	4	15.2	76.0	40	ω	38-466-588-5	52, A
	S	15	٦.	15	-	00-1	37, A
	ωı	12	٦.	15	17	-08-377-310-3	37,
	7	12	75.0	15		-09-757-100B-	m
	20 (14.8	٠.	35		144-428-	31,
		14.8		35		1 (m (
	0 :	14.8	74.0	3.5	8 0	US-09-441-966-31	Sequence 31,
	- C	* *		, ,	٥ -	US-US-4/2-801-1314	1014
	٦,	# C	0.77	000	2 -	US-U8-666-233-1314	Sequence 1314, Ap
ر	+ -	7.4.7		2 6		FCI - 033/ - 00104 - 10	٠,
C	15	13.6	0.00	25	5.5	US-60-234-017-326383	32638
טיי	٦,	13.4		18	14	US-09-082-614-16	16. A
O	1	13.2		19		US-60-216-745-5602	5602
υ	7	13.2		20	13	US-08-965-620-718	718, P
υ	1	13.2	0.99	25	26	-660-220-1	13689
O	~	ж Э	٠.	25	52	-60-233-166-5	57027
υ	C)	m,	•	25	55	-60-233	384424
O	(1)	۳,	٠,	43	1,3	-310-298-3092	3092,
υ	N (٦,		25	5	-166-2	260192,
O		٠		19	77	US-09-696-791-539	539,
	070	27.0	0.4	200	9 0	US-US-/US-/US-TT3US	Sequence 11309, A
	0 6			0 0	9 0	US-03-703-700-17070 US-60-164-320-11309	11300,
	28	12.8	64.0	20	8 4	US-60-164-320-17076	17076.
	29			20	50	US-60-183-791-11309	11309,
	30		64.0	20	20	8-60-183-791-1707	H
	31		•	21	⊣ '	17-07972-3	Sequence 3, Appli
	32			21	10	647-351A-	m
		٠		23	5:	100-649A	m (
	3.4 2.0	•	•	77	1	,	Sequence 3, Appli
	י ר	•		22	0 4	733-100-2124	Sequence 22124U,
יכ				200	ט מ	234-01/-90	Sequence 9000, Ap
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טנ	יי רי		* =	2 C	U L	-234-01/-3364	
) (7		* <	2 0	י ני	-60-234-017-33645	33645
י כ	4			3.5	ς α	-09-465-684-134	134. AD
	•		4	41	18	-09-404-520-30	30598
O	4		c	20	7	PCT-US00-00583-43	<u></u>
	44	12.6	63.0	25	55	-60-233-166-70	07
O	45	12.6	63.0		22	US-60-234-017-229508	Sequence 229508,

ALIGNMENTS

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LENGTH: 40 base pairs
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STRANDEDNESS: single
TOPOLOGY: linear
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC
OPERATING SYSTEM:
SOFTWARE: Patent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: Illinois
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 321 NC
CITY: Chicago
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US-08-466-588-52/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
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                                                                                                                                                                                                                         APPLICANT: Nero, Pamela S.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase.
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310A
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 17
LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 20; DB 1;
Pred. No. 33;
                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: PCT/USO0/18999; CURRENT FILING DATE: 2000-07-13; PRIOR APPLICATION NUMBER: 09/377,310 PRIOR FILING DATE: 1999-08-19; NUMBER OF SEQ ID NOS: 43 SOFTWARE: Patentin Ver. 2.0 SEQ ID NO 1; SE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                      Isis Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: antisense sequence PCT-US00-18999-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; OTHER INFORMATION: antisense sequence US-09-377-310-17
                                   Sequence 17, Application PC/TUS0018999 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17, Application US/09377310A GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17, Application US/09757100B
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100.0%; Pr
                                                                                               1 attectegetgetggaa 20
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Best Local Similarity 100.
Matches 20; Conservative
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Best Local Similarity 100.
Matches 20; Conservative
PCT-US00-18999-17
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US-09-757-100B-17
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Gaps
CURRENT HOUSE, Brett P.

APPLICANT: Monia, Brett P.

APPLICANT: Gaarde, William A.

APPLICANT: Garde, William A.

APPLICANT: Garde, William A.

FIGURENT: Nero, Pamela S.

TITLE OF INVENTION: Expression

FILE REFERENCE: ISPH-0533

CURRENT FILING DATE: 2001-03-15

PRIOR FILING DATE: 1999-08-19

PRIOR FILING DATE: 1999-08-19

PRIOR FILING DATE: 2000-07-13

NUMBER OF SEQ ID NOS: 44

SOFTWARE: Patentin Ver: 2.0

SEQ ID NO 17

LENGTH: 20

FILENCE: A CONTROL OF THE CONTROL 
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TITLE OF INVENTION: METHODS AND MATERIALS RELATING TO THE
TITLE OF INVENTION: FUNCTIONAL DOMAINS OF DNA BINDING PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;
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Pred. No. 33;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Arnold, White & Durkee
STREET: 321 North Clark Street, Suite 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             red. No. 33;
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
ATORNEY, AGENT INFORMATION:
NAME: COUGHLIN, Daniel F.
REGISTRATION NUMBER: 36,111
REFERENCE/DOCKET NUMBER: arcd067
TELECHONE: (312) 744-0090
TELEPHONE: (312) 744-0091
INFORMATION FOR SEQ ID NO: 52:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CTHER INFORMATION: antisense sequence US-09-757-1008-17
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APPLICATION NUMBER: US/08/466,588
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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; GENERAL INFORMATION:
APPLICANT: Sukhatme, Vikas P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/040,548
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
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Best Local Similarity 100.0%;
Matches 20; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
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Probe oligo-1 for Probe used to isol Probe used to isol Probe used to isol

for human J Immunoglobul heavy chain

Human

AAV12514 AAT70497 AAT73468 AAT73448

Нитап

J-mu region

Newcastle disease Newcastle disease Primer ELO74 for N PCR primer ELO74 u Human MDC PCR prim

AAV42526 AAQ76135

PCR primer SEQ ID Delta-9 desaturase

thrombopoiet flt1 VEGF re Hammerhead ribozym Human Mer receptor Human Ig H chain s Human Ig H chain s Rat ICAM hammerhea

Human Human

AAA99419 AAX63148 AAZ87393 AAX68545 AAF00661 AAT53885 AAT40112 AAA78278

Oligonucleotide SE Primer 1RAF616 for FAR1 gene amplific

AAV38155 AAV38128 AAZ22071 AAZ22071 AAX06008 AAY82530 AAV82530 AAV85543 AAX65443 AAX65443 AAX65443 AAX3524 AAT3524 AAT35524

Mouse B7-1 hammerh Primer EL074 to ge

US5874299 Seq ID 1 Apolipoprotein A g Probe Aposma-1 use

Rat farnesyl prote JH probe #3 to det Antifreeze protein Synthetic antifree

AAA78291 AAQ94416 AAQ30098

AAX88102 AAX88103

ALIGNMENTS

Run

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New antisense compounds for inhibiting focal adhesion kinase expression, especially useful for inhibiting retinal neovascularization, or for diagnosing and treating e.g. colon cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; focal adhesion kinase; FAK; signal transduction; cancer; embryonic development disorder; anglogenic disorder; wound healing; antisense; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human focal adhesion kinase antisense sequence #10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP.
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 AAC65544 standard; DNA; 20
  (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaarde WA;
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Homo sapiens.
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Primer mcl-like-1r
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14.320 Million cell updates/sec
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(c) 1993 - 2000 Compugen Ltd.
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                                    The present invention describes a number of phosphorothioate antisense sequences to the human focal adhesion Kinase (FAK) protein. This protein is involved in integrin-mediated signal transduction, and is implicated in cancer, particularly colon, breast and oral tumours, embryonic development disorders, angiogenic disorders and wound healing. The antisense sequences, including the one shown here, can be used in the treatment of all of these.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; BRG1; tumour suppressor gene; cancer; chromosome 19p13.1;
retinoblastoma tumour suppressor gene; RB; drug screening; gene therapy;
drug design; peptide therapy; animal model; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Diagnosing a polymorphism associated with predisposition for cancer in humans by determining whether there is a germline alteration of a BRG1 gene or its expression products -
                                                                                                                                                                                                                                                                                                                  Gaps
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Claim 15; Column 23; 30pp; English
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Matches 20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compounds for inhibiting focal adhesion kinase expression, especially useful for inhibiting retinal neovascularization, or for diagnosing and treating e.g. colon cancer
                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                    Human; focal adhesion kinase; FAK; signal transduction; cancer; embryonic development disorder; angiogenic disorder; wound healing; antisense; phosphorothioate; ss.
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    Length 20;
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100.0%; Pred. No. 1.1e+02;
iive 0; Mismatches 0;
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Pred. No. 22;
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94.4%;
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                                              Conservative
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Best Local Similarity 100.
Matches 15, Conservative
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Matches 17; Conserv
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MC-R1; melanocortin 1 receptor; rhodopsin; G-protein coupled receptor; splice variant; MC-R1B; C-terminal extension; agonist; antagonist; modulator; PCR primer; ss.
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                                                                                                                                                                                                                                 (MERI ) MERCK & CO INC.
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Best Local Similarity
Matches 16; Conserv
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                                                                  Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention provides a method of diagnosing a vascular disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       In an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also useful in forensics, paternity testing, genetic analysis and phenotype correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification.
                                              Human; variant thrombospondin 1; variant thrombospondin 4; SNP;
polymorphism; vascular disease; coronary artery disease; forensics;
myocardial infarction; atherosclerosis; stroke; venous thromboembolism;
pulmonary embolism; paternity test; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                McCarthy JJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and atherosclerosis
                                                                                                                                                                                                /*tag= a
/standard_name= "single nucleotide polymorphism"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Primer mc1-like-2f for human MC-R1 exon 2 amplification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Daley GQ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14.2; DB 22;
Pred. No. 2.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .;
                 Human gene single nucleotide polymorphism #253.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bolk S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21 BP; 7 A; 4 C; 7 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                             WHED ) WHITEHEAD INST BIOMEDICAL RES.
                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ireland JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Examples; Page 67; 242pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
                                                                                                                                                                                 replace(11, A)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         71.0%;
84.2%;
                                                                                                                                                                                                                                                                                                                                              99US-0153357.
                                                                                                                                                                                                                                                                                                                                                                                                                              (MILL-) MILLENNIUM PHARM INC
                                                                                                                                                                                                                                                                                                                                                           26-JUL-2000; 2000US-0220947.
                                                                                                                                                                                                                                                                                                             07-SEP-2000; 2000WO-US24503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cctgacatcagtagcatct 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19 CTTGCCATCAGTGGCATCT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA51849 standard; DNA; 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-JAN-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 71.03
Best Local Similarity 84.23
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gargill M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-226749/23
                                                                                                                                                                                                                                              WO200118250-A2
                                                                                                                                                                                                                                                                                                                                          10-SEP-1999;
                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                               15-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lander ES,
                                                                                                                                                                               Variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA51849;
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RESULT

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The invention concerns novel splice variants of the melanocortin 1 receptor (MC-R1) protein belonging to the rhodopsin sub-family of G-protein coupled receptors. The splice variants, referred to as MC-R1B proteins, contain an intracellular domain with an additional 65 amino acid residues in comparison to previously disclosed human MC-R1, referred to as MC-R1B, additionally, residue 317 of the MC-R1B proteins is Cys, whereas the C-terminal amino acid residue 317 of the MC-R1B proteins is Cys. Trp. The novel sequences can be used to determine whether a substance antagonists of MC-R1B. Pharmaceutical compositions comprising modulators of MC-R1B are used to treat or diagnose disorders involving
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MC-R1; melanocortin 1 receptor; rhodopsin; G-protein coupled receptor; splice variant; MC-R1B; C-terminal extension; agonist; antagonist; modulator; PCR primer; ss.
                                                                                                                                                                   New nucleic acid encoding a human melanocortin 1 receptor protein (MC-R1B) for determining whether a substance is capable of binding to or activating human MC-R1B and identifying a substance that modulates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Primer mc1-like-lr for human MC-R1 exon 1 amplification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             inappropriate melanocortin expression or activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                68.0%; Score 13.6; DB 21;
1larity 80.0%; Pred. No. 6.4e+02;
Conservative 0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 39 BP; 5 A; 16 C; 9 G; 9 T; 0 other;
Van Der Ploeg LHT;
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Example 14; Page 82; 172pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-MAR-1992;
23-JUN-1992;
                                                                                                                                                                                                                                                                                                      10-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                  WO9312227-A.
                                                                                                                                                                                                                                                                                                                                                                                                                      24 - JUN-1993
                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                  AAQ44160;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kay RM,
                                                                                                                                                                                                                                                    AAQ44160/c
                                                                                                                                                                                                                                          RESULT
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                                                                                                                                                                          receptor (MC-R1) protein belonging to the rhodopsin sub-family of expotein coupled receptors. The splice variants, referred to as MC-R1B proteins, contain an intracellular domain with an additional 65 amino acid residues in comparison to previously disclosed human MC-R1, referred to as MC-R1A. Additionally, residue 317 of the MC-R1B proteins is Cys, whereas the C-terminal amino acid residue 317 of known MC-R1B proteins is Trp. The novel sequences can be used to determine whether a substance antagonists of MC-R1B receptor activity and to identify potential agonists or modulators of MC-R1B are used to treat or diagnose disorders involving inappropriate melanocortin expression or activity.
                                                                                          New nucleic acid encoding a human melanocortin 1 receptor protein (MC-RIB) for determining whether a substance is capable of binding to or activating human MC-RIB and identifying a substance that modulates MC-RIB receptor activity
                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                  The invention concerns novel splice variants of the melanocortin 1
                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                     Length 40;
                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Immunoglobulin trans:genes - for prodn. of heterologous non-rearranged and/or rearranged Ig chains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human heavy chain J region specific oligonucleotide.
                                                                                                                                                                                                                                                                                                                                   Score 13.6; DB 21;
Pred. No. 6.4e+02;
0; Mismatches 4;
                                                                                                                                                                                                                                                                                                     Sequence 40 BP; 8 A; 12 C; 13 G; 7 T; 0 other;
                                                  Van Der Ploeg LHT;
                                                                                                                                              Example 1; Page 49; 101pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Placental; probe; heavy chain; ss.
                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP
                                                                                                                                                                                                                                                                                                                                    68.0%;
80.0%;
                                                                                                                                                                                                                                                                                                                                                                          1 cctgacatcagtagcatctc 20
|||||| | ||| |||| |||
9 cctgagagcaggagcatgtc 28
         98US-0113401
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAQ23448 standard; DNA; 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-AUG-1992 (first entry)
                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 80.0
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GENP-) GENPHARM INT INC.
                                                  MacNeil DJ,
                              (MERI ) MERCK & CO INC
                                                                     WPI; 2000-452365/39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1992-113962/14.
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        23-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-AUG-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          31-AUG-1990;
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                                                  Howard AD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9203918-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-MAR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lonberg N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAQ23448;
                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ23448/C
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A human placental genomic DNA library cloned into the phage vector lambda EMBL3/SP6/T7 was screened with the human heavy chain J region specific oligonucleotide probe and the phage clone lambda 1.3 isolated. A 6 kb HindII/KpuI fragment from this clone, contg. all six J segments as well as D segment DHQ22 and the heavy chain J-mu intronic enhancer was isolated, and ultimately used to construct an IGM expressing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A 6.3kb BamHI/HindIII fragment that includes all J segments was isolated from human genomic DNA library using probe AAQ44160. An adjacent 10kb HindIII/BamHI fragment that contains enhancer, switch and constant region coding exons was similarly isolated using the probe AAQ44161 and an adjacent 3' 1.5kb BamHI fragment was isolated using pMUM insert as probe and cloned into pUCl9. (pMUM is a 4kb EcoRI/HindIII fragment isolated from human genomic DNA library with oligonucleotide AAQ44162 mu membrane exon 1). All three fragments are used in the construction of plasmid pHIG1 which contains an 18kb insert encoding J and C-mu segments.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Immunoglobulin; Ig; heavy chain; minilocus; isotype switching; J region; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                     Length 42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Probe to isolate human immunoglobulin Joining segments
                                                                                                                                                                                                                                                                                                                                                                                                                  68.0%; Score 13.6; DB 13;
80.0%; Pred. No. 6.4e+02;
iive 0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                 Sequence 42 BP; 5 A; 8 C; 15 G; 14 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 42 BP; 5 A; 8 C; 15 G; 14 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 4; Page 60; 196pp; English.
                                                                                                                                                                                                                                             See also 023419-50, 022417-30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 cctgacatcagtagcatctc 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     35 CCAGACATCAAAAGCATCAC 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               92WO-US10983.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GENP-) GENPHARM INT INC
                                                                                                                                                                                                         mini-locus transgene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1993-214169/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Lonberg N;
                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 16; Conserv
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(first entry)

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Heavy chain; J region; human; probe; minilocus; transgene; transgenic; mouse; production; heterologous; antibody; IgM; immunoglobulin; ss.
                                                                                                                   Human antibody heavy chain J region specific probe.
AAT37238/c
ID AAT37238 standard; DNA; 42 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GENP-) GENPHARM INT INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1996-383736/38.
                                                                                                                                                                                                                                                                                                                                                                    29-AUG-1990;
31-AUG-1990;
17-DEC-1991;
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                                                                                    18-APR-1997
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                                                                                                                                                                                                                                                                                   13-AUG-1996
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                                                                                                                                                                                                                     Synthetic.
                                                   AAT37238;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kay RM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 11
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                                                   ó
                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A human placental genomic DNA library cloned into the phage vector lambda EMBL3/SP6/T7 was screened with the human heavy chain J-region specific oilgonucleotide 044180. Phage clone lambda 1.3 was isolated. A 6kb HindIII/KpnI fragment from this clone, containing all 6 J segments as well as D segment DH022 and the heavy chain J-mu intronic enhancer, was isolated. The same library was screened with the human mu specific oligonucleotide 044181 and phage clone mu switch region and all of the mu constant region as and all of the mu constant region as solated from this clone. The two fragments were ligated together
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  th KpnI/XhoI digested pNNO3 to obtain plasmid pJMI which was used the construction of IgM expressing minilocus transgene pIGM1.
                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Transgenic non-human animals contg. immunoglobulin heavy chain trans gene - used to produce useful antibodies by isotype switching
               68.0%; Score 13.6; DB 14; Length 42; ilarity 80.0%; Pred. No. 6.4e+02; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  68.0%; Score 13.6; DB 14; Length 42; 80.0%; Pred. No. 6.4e+02; 1ve 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                   Immunoglobulin; IgM; heavy chain; minilocus transgene;
isotype switching; J-mu constant region; Joining region; ss.
                                                                                                                                                                                                                                                                                                 Probe specific for human heavy chain J region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 42 BP; 5 A; 8 C; 15 G; 14 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 12; Page 91; 196pp; English.
                                                                                                                                                                                                  AAQ44180 standard; DNA; 42 BP.
                                                                                  20
                                                                                                   35 CCAGACATCAAAAGCATCAC 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   91US-0810279.
92US-0853408.
92US-0904068.
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                                                                                                                                                                                                                                                                    (first entry)
                                                                                  1 cctgacatcagtagcatctc
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                                  Local Similarity
les 16; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-DEC-1992;
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                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                   AAQ44180;
                    Query Match
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                                  Best Loca
Matches
                                                                                                                                                                                 AAQ44180/c
                                                                                                                                                                 RESULT
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90US-0574748. 90US-0575962. 91US-0810279.

92US-0853408 92US-0904068

Lonberg N;

90US-0574748 92US-0990860

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                      chain J region, which was used in the construction of a IgM heavy chain minilocus transgene. The transgene was injected into mouse embryo pronuclei to generate transgenic mice, which can be used for the production of heterologous (i.e. human) antibodies against specific antigens, this comprises immunising a mouse with a preselected antigen and collecting antigen binding heterologous
                                                                                                                                           The present sequence is a probe specific for a human antibody heavy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;
0
of heterologous human immunoglobulin(s) - by immunising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Heavy chain; gene segment; human; DNA fragment; probe; minilocus; transgene; transgenic; mouse; J segment; production; heterologous; antibody; gamma; immunoglobulin; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              68.0%; Score 13.6; DB 17; 80.0%; Pred. No. 6.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human antibody J segment oligonucleotide probe.
                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 42 BP; 5 A; 8 C; 15 G; 14 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                 Example 12; Columns 57-58; 94pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;
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ID AAT37216 standard; DNA; 42 BP
XX
XX
XX
XX
DT 18-APR-1997 (first entry)
XX
DE Human antibody J segment olig
XX
XX
DE Human intibody J segment, hum
XX
XX
W Heavy chain; gene segment; hum
XX
XX
W Hooduction; heterologous; ant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    35 CCAGACATCAAAAGCATCAC 16
                                                                                                                                                                                                                                                                                                                                                  human IgM immunoglobulins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 cctgacatcagtagcatctc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 16; Conserv
                           transgenic mice
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Gaps

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ò g 10

RESULT

16; Conservative

Local Similarity

Best Loca Matches

us-09-757-100b-12.szlim50.rng

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Human antibody producing transgenic mouse – containing transgene comprising human V, D and J genes and sequences to provide isotype
                                                                                                                                                                                                                                                                                                         Example 4; Column 50; 153pp; English.
                                                                                     92US-0853408.
92US-0904068.
92US-0990860.
                                                                                                                               93US-0096762.
93US-0155301.
93US-0161739.
                                            90US-0574748
90US-0575962
            94US-0352322
                                                                 91US-0810279
                                                                           92US-0834539
                                                                                                                      93US-0053131
                                                                                                                                                                 93US-0165699
                                                                                                                                                                          94US-0209741
                                                                                                                                                                                                (GENP-) GENPHARM INT INC.
                                                                                                                                                                                                                                                                                      switching in lymphocytes
                                                                                                                                                                                                                                         WPI; 1997-258277/23.
                                                                                                                                                                                                                     Lonberg N;
           07-DEC-1994;
                                                                                                                                 22-JUL-1993;
                                                                                                                                            .8-NOV-1993;
                                                                                                                                                     03-DEC-1993;
                                                                                                                                                                 .0-DEC-1993;
                                                                                       18-MAR-1992
                                                                                                            16-DEC-1992
                                                                                                                      26-APR-1993
                                                                 17-DEC-1991
                                                                           05-FEB-1992
                                                                                                 23-JUN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAT70497;
                                                                                                                                                                                                                    Кау км,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAT70497/C
ID AAT704
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 13
  Dp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Transgenic mouse; human; immunoglobulin; heavy chain segment; J region; joining region; constant region; VH family; variable gene; gamma isotype; diversity gene; isotype switching sequence; mu isotype; Ig production; monoclonal antibody; MAb production; antigen; heavy chain isotype;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                     BamHI/Hindill fragment that includes all human antibody J segments. The fragment isolated using the probe was used in the construction of a heavy chain minilocus transgene. The transgene, which contains human immunoglobulin heavy chain coding sequences, can be injected into mouse embryo pronuclei to generate transgenic mice. The mice can be used for the production of heterologous (i.e. human) antibodies against specific antigens, this comparises immunising a mouse with a preselected antigen and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
                                                                                                                                                                                                                                       of heterologous human immunoglobulin(s) - by immunising
                                                                                                                                                                                                                                                                                            present sequence is an oligonuclectide probe for a 6.3 kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                         68.0%; Score 13.6; DB 17; Length 42; llarity 80.0%; Pred. No. 6.4e+02; Conservative 0; Mismatches 4; Indels (
                                                                                                                                                                                                                                                                                                                                                                                           collecting antigen binding heterologous human gamma
                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 42 BP; 5 A; 8 C; 15 G; 14 T; 0 other;
                                                                                                                                                                                                                                                                        Example 4; Column 36; 94pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Probe for human J region sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antigenic stimulation; probe; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAV12514/c
ID AAV12514 standard; DNA; 42 BP
                                                                                                          90US-0574748.
90US-0575962.
91US-0810279.
92US-0853408.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cctgacatcagtagcatctc 20
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                                                                          90US-0574748
                                                                                                                                                     92US-0904068
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-MAY-1998 (first entry)
                                                                                                                                                                        (GENP-) GENPHARM INT INC.
                                                                                                                                                                                                                   WPI; 1996-383736/38.
                                                                                                                                                                                               Kay RM, Lonberg N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                    transgenic mice
                                                                          29-AUG-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                               US5545806-A
                                                                                                                     31-AUG-1990;
17-DEC-1991;
                                                     13-AUG-1996
                                                                                                16-DEC-1992
                                                                                                          29-AUG-1990
                                                                                                                                         18-MAR-1992
                                                                                                                                                     23-JUN-1992
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          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
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                                                                                                                                                                                                                                       Prodn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 12
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This sequence represents a probe for the human junction (J) region genes. The identified sequence is used in the plasmid pRE3, which is used to develop the transgenic mouse of the invention contains in its genome a transgene comprising in mouse of the invention contains in its genome a transgene comprising in operable linkage human variable (V), diversity (D) and J genes, a human of mu constant region gene (muCH), at least 2 different non-mu human CH genes and associated isotype switching sequences, where human mu and gamma switch sequences are located in closer proximity to each other than in the naturally occurring human immunoglobulin (Ig) locus, and where in lymphocytes of the mouse the transgene undergoes productive VDJ rearrangement and mu to gamma isotype switching by recombination between the human mu and gamma sequences, so that the mouse produces a serum containing Ig of at least 3 human hawy chain isotypes in response to antigenic stimulation. The transgenic mice can be used to produce human contigens. The MAD can be used in therapeutic or diagnostic human antigens. The transgenic mice can produce human MAD of multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      68.0%; Score 13.6; DB 18; Length 42; 80.0%; Pred. No. 6.4e+02; Live 0; Mismatches 4; Indels (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human Immunoglobulin J region oligonucleotide probe 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 42 BP; 5 A; 8 C; 15 G; 14 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                isotypes by undergoing isotype switching.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT70497 standard; DNA; 42 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 cctgacatcagtagcatctc 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             35 CCAGACATCAAAAGCATCAC 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-MAR-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 68.0
Best Local Similarity 80.0
Matches 16; Conservative
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A novel composition has been developed which comprises an immunoglobulin (Ig) having an affinity constant (Ra) of at least 2 multiply 10000000000 M-1 for binding to a predetermined human antigen. The 1000000000 M-1 for binding to a predetermined human antigen. The present sequence represents a probe used to screen a human placental genomic DNA library cloned into the phage vector lambdaEMBL3/SP6/T7, for the human heavy chain J region. Anti-CD4 antibodies may be used in therapeutic and diagnostic applications, especially for the treatment of human diseases. These antibodies reduce activity of CD4 cells and transplant rejection. Transgence animals are capable of producing heterologous antibodies of multiple isotypes by undergoing isotype switching. These animals produce a first ig type that is necessary for antigen-stimulated B-cell maturation and can switch to encode and produce one or more subsequent heterologous isotypes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A novel composition has been developed which comprises an immunoglobulin (Ig) having an affinity constant (Ka) of at least 2 multiply
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ig; affinity constant; human; antigen; hybridoma; B cell; transgene; transgenic; mouse; CD4; antibody; autoimmune; inflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel anti-CD4 antibody produced by transgenic mice - used in the treatment of auto-immune disease etc.
                                                                             Novel anti-CD4 antibody produced by transgenic mice - used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;
0
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80.0%; Pred. No. 6.4e+02;
ive 0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 42 BP; 5 A; 8 C; 15 G; 14 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human J-mu region genomic DNA library probe.
                                                                                                  treatment of auto-immune disease etc.
                                                                                                                                   Example 12; Page 133; 396pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 4; Page 103; 396pp; English.
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ID AAT73448 standard; DNA; 42 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 cctgacatcagtagcatctc
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Best Local Similarity 80.05
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      transplant rejection; ss.
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                                     WPI; 1997-235888/21
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Kay RM, Lonberg N;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-OCT-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This oligonucleotide was used as a probe to facilitate in the cloning of a J-mu region from a human immunoglobulin. The probe isolated a 6.3 kb BamH/HindIII fragment from a human genomic library which included all joining (J) segments. This region is used in a novel method of developing transgenic non-human animals capable of producing heterologous antibodies encoded by human immunoglobulin genes. Such transgenically produced monoclonal antibodies should alleviate the intrinsic immunogenicity of non-human immunoglobulins allowing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ig; affinity constant; human; antigen; hybridoma; B cell; transgene; transgenic; mouse; CD4; antibody; autoimmune; inflammatory; transplant rejection; immunoglobulin; ss.
                                                                                                                                                                                                                                                                                                                                                                               Transgenic mouse for heterologous antibody production - containing DNA encoding human immunoglobulin components
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Pred. No. 6.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 42 BP; 5 A; 8 C; 15 G; 14 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human heavy chain J region specific probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            development of new in vivo applications.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 5; Column 33; 90pp; English.
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ID AAT73468 standard; DNA; 42 BP.
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80.0%;
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                                                                                                                                                          92US-0834539
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                                                                                                                                                                                                                                                        GENP-) GENPHARM INT INC.
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                                                                                                                                                                                                                                                                                                Lonberg N;
                                                                                                                                                                                                                                                                                                                                        WPI; 1997-297410/27
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Best Local Similarity
                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
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                                                                                                                                                        05-FEB-1992;
                                                                                                                                                                                              29-AUG-1990;
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                                                                         US5633425-A.
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                                                                                                                27-MAY-1997
                Synthetic
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1000000000 M-1 for binding to a predetermined human antigen. The present sequence represents a probe used for the isolation of human J segments from a human genomic DNA library. The anti-CD4 entibodies may be used in therapeutic and diagnostic applications, especially for the treatment of human diseases. These antibodies reduce activity of CD4 cells and reduce undesirable autoimmune reactions, inflammatory response and transplant rejection. Transgenic animals are capable of producing heterologous antibodies of multiple isotypes by undergoing isotype switching. These animals produce a first Ig type that is necessary for antigen-stimulated B-cell maturation and can switch to encode and produce one or more subsequent heterologous isotypes.
             88888888888888888888888888888888888888
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Sequence 42 BP; 5 A; 8 C; 15 G; 14 T; 0 other;

; Gaps ö Query Match 68.0%; Score 13.6; DB 18; Length 42; Best Local Similarity 80.0%; Pred. No. 6.4e+02; Matches 16; Conservative 0; Mismatches 4; Indels (

οy g

Search completed: October 2, 2001, 16:18:41 Job time: 15485 sec

Sequence Seq

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Sequence 11, Application US/09377310B

Patent No. 6133031

GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT FILING DATE: 1999-08-19
CURRENT FILING DATE: 1999-08-19
SOFTWARE: EatentIn Ver. 2.0
SEQ ID NO.11

SEQ ID NO.11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 31, Application US/09377310B
Patent No. 6133031
GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase: TITLE OF INVENTION: Expression
FILLE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310B
CURRENT APPLICATION NUMBER: US/09/377,310B
NUMBER OF SEQ ID NOS: 1999-08-19
SOFTWARE: PatentIn Ver. 2.0
US-08-483-389-87
US-08-483-513D-87
US-08-473-503-87
US-08-473-503-87
US-08-720-420A-87
US-08-720-420A-87
US-08-475-680-87
US-08-483-389-100
US-08-783-503-100
US-08-783-503-100
US-08-783-503-106
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100.0%; Pred. No. 0.034;
.ive 0; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                     ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: antisense sequence US-09-377-310-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 agtacccaggtgagtcttag 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 agtacccaggtgagtcttag 20
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20; Conservative
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 SEQ ID NO 31
LENGTH: 15
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Best Local S
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Sequence 8, Appli
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Description
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/PcTUS_COMB.seq:*
                 GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-377-310-31
US-08-136-118-7
US-08-136-118-7
US-08-530-492-69
US-08-906-517-69
US-08-908-128-15
US-09-237-510-6
US-09-137-510-6
US-09-117-952-668
US-09-117-952-668
US-09-06-410-12
US-08-685-871-14
US-08-685-871-33
US-08-685-871-33
US-08-685-871-41
US-08-685-871-41
US-08-685-871-43
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US-08-685-871-43
US-08-685-871-43
US-08-685-871-43
US-08-685-871-43
US-09-993-4
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US-08-818-604-8
US-08-482-882-87
                                                                                                                                                                                                                                                                                                                               324599 seqs, 94655562 residues
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Maximum Match 100%
Listing first 45 summaries
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score greater than
and is derived by
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Length 20; 0; Indels

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RESULT 5
US-08-530-492-69/c
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APPLICANT: HOKE, Matthews O
APPLICANT: BRADLEY, Matthews O
APPLICANT: WILLIAMS, Taffy J
APPLICANT: LEE, Che-Hung
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDES DIRECTED
TITLE OF INVENTION: AGAINST HUMAN ICAM-1
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
ADDRESSEE: Naval Medical Res. & Dev. Cmd.
STREET: Bethesda
STATE: Maryland
CUTY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 13.4; DB 1; Length 21;
Pred. No. 1.2e+02;
0; Mismatches 1; Indels
                                                                                  DB 3; Length 15;
16;
                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/136,118
                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 07/918,259
FILING DATE: 24-UUL-1992
ATTORNEY/AGENT INFORMATION:
NAME: Spewack, A. David
REGISTRATION NUMBER: 24,743
REFERENCE/DOCKET NUMBER: N.C. 75,776
TELECOMMUNICATION INFORMATION:
                                                                                  75.0%; Score 15;
100.0%; Pred. No.
ive 0; Mismatc
       ; OTHER INFORMATION: antisense sequence US-09-377-310-31
                                                                                                                                                                                                                                                                          Sequence 7, Application US/08136118
Patent No. 5580969
GENERAL INFORMATION:
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TELEFRAX: (202) 295-1022
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               67.0%;
93.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 20889-5606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                  Query Match 75.0
Best Local Similarity 100.
Matches 15; Conservative
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TYPE: nucleic acid
STRANDEDNESS: single
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Best Local Similarity
Matches 14; Conserv
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FEATURE:
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Parent No. 5689052
GENERAL INFORMATION:
APPLICANT: Brown, Sherri M.
APPLICANT: Fromm, Michael E.
APPLICANT: Sanders, Particia R.
APPLICANT: Sanders, Part
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                                                                                                                                APPLICANT: Yeh, Homer R., Wick, Charles H.
TITLE OF INVENTION: NOVEL DETECTOR FOR NUCLEIC ACID
TYPING AND METHODS OF USING THE SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      64.0%; Score 12.8; DB 4; Length 27; 70.0%; Pred. No. 2.5e+02; Live 2; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: Maryland
COUNTRY: United States of America
ZIP: 21010-5423
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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700 Chesterfield Parkway No. 5689052th
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CHROMOSOME/SEGMENT: EFD52 Minisatellite;
SEQUENCE DESCRIPTION: SEQ ID NO: 8:
US-09-246-277A-8
                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 13
CORRESPONDENCE ADDRESS:
ADDRESSEE: Office of the Chief Counsel
STREET: U.S. Army SBCCOM
CITY: APG(EA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/09/246,277A FILING DATE: 08-Feb-1999 CLASSIFICATION: <Unknown>
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NAME: Biffoni, U. John
REGISTRATION NUMBER: 39908
REFERENCE/DOCKET NUMBER: 436-96
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS: not relevant
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Sequence 8, Application US/09246277A Patent No. 6238866 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (410) 436-2534
INFORMATION FOR SEQ ID NO: 8:
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MOLECULE TYPE: DNA (genomic)
HYPOTHETICAL: NO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 27 base pairs
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Best Local Similarity 70.0
Matches 14; Conservative
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ORIGINAL SOURCE:
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                                                                                                                                                                                                                                                                          Length 29;
                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Ting, Jenny Pan-Yung
APPLICANT: Chin, Keh-Chin
TITLE OF INVENTION: No. 5994505el Forms of Class II MHC
TITLE OF INVENTION: Transactivator (CIITA)
UNMBER OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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COMPUTER READABLE FORM:
COMPUTER: TEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NATA:
APPLICATION NUMBER: US/08/988,128
FILING DATE: 10-DEC-1997
                                                                                                                                                                                                                                                                        Score 12.8; DB 4;
Pred. No. 2.5e+02;
0; Mismatches 2;
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Pred. No. 2.6e+02;
0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 7
US-08-988-128-15
Sequence 15, Application US/08988128
Patent No. 5994505
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Raleigh.
STATE: No. 5994505th Carolina
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Biswas, Sorojini J.
REGISTRATION NUMBER: 39,111
REFERENCE/POCKET NUMBER: 5470
TELECOMMUNICATION INFORMATION:
TELEFAN: (919) 854-1400
TELEFAN: (919) 854-1401
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
REGISTRATION NUMBER: 33,928
REFERENCE/DOCKET NUMBER: MOB
TELECOMMUNICATION INFORMATION:
TELEPHONE: 512-418-3000
TELEFAX: 512-418-377
INFORMATION FOR SEQ ID NO: 69: SEQUENCE CHARACTERISTICS:
LENGHH: 29 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; MOLECULE TYPE: DNA (genomic) US-08-988-128-15
                                                                                                                                                                                                                                                                        64.08;
87.58;
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87.5%;
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Best Local Similarity 87.5
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3 tacccaggtgagtct 18
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16 TACCCAGCTGTGTCTT 31
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                                                                                                                                                                                                                                                                                                                                                                            24 AGTGCCCAGGAGAGTC 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nucleic acid
EDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 14; Conserva
                                                                                                                                                                                               linear
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                                                                                                                                                                                               ; TOPOLOGY:
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APPLICANT: Brown, Duff A.
APPLICANT: Dean, Duff A.
APPLICANT: Dean, Michael B.
APPLICANT: Sanders, Partricia R.
TITLE OF INVENTION: Expression in Monocotyledonous Plants and Method For TITLE OF INVENTION: Preparation Thereof CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
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Pred. No. 2.5e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/906,517
FILING DATE: COGGUITZENTY HEREWITH
                                                                                                                                                    SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/530,492
                                                                                                                                                                                                         FILING DATE:
CLASSIFICATION: 800
PELORATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/08/172,333
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: HOERING T, Dennis R.
REGISTRATION NUMBER: 30,914
REFERENCE/DOCKET NUMBER: 38-21(10605)A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (314537-6099
INFORMATION FOR SEQ ID NO: 69: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Arnold, White & Durkee
STREET: P.O. Box 4433
CITY: Houston
                                                                                                               COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 69, Application US/08906517
Patent No. 6180774
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; MOLECULE TYPE: DNA (synthetic) US-08-530-492-69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kitchell, Barbara S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 87.5%;
Matches 14; Conservative
                                                                                                 Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 agtacccaggtgagtc 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 29 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24 AGTGCCCAGGAGAGTC 9
                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
St. Louis
Missouri
                                          USA
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                                                           63198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
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                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: F
                    STATE:
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Gaps

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APPLICANT: Robert McKay
APPLICANT: Madeline M. Butler
APPLICANT: Madeline M. Butler
APPLICANT: Jacqueline Wyatt
APPLICANT: Jacqueline Wyatt
APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF PEPCK-CYTOSOLIC EXPRESSION
TITLE OF INVENTION: ANTISENSE MODULATION OF PEPCK-CYTOSOLIC EXPRESSION
CURRENT APPLICATION NUMBER: US/09/488,671A
CURRENT FILING DATE: 2000-01-21
SEQ ID NO 119
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                Length 48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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Patent No. 5851760
GENERAL INFORMATION:
APPLICANT: Evans, Glen A.
APPLICANT: Smith, Michael W.
TITLE OF INVENTION: SAMPLED MAPS OF COMPLEX GENOMES
UNDMBER OF SEQUENCES: 797
CORRESPONDENCE ADDRESS:
ADDRESSE: Pretty, Schroeder, Brueggemann & Clark
                                       SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/237,510
                                                                                                                                                                                                                                                                                                                                                                                                                           Score 12.8; DB 2;
Pred. No. 2.7e+02;
); Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: Antisense Oligonucleotide US-09-488-671-119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; Sequence 119, Application US/09488671A; Patent No. 6187545; GENERAL INFORMATION:
COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                      CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: FUGIt, Donna R.
REGISTRATION NUMBER: 32,135
REFERENCE/DOCKET NUMBER: P-37,
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                           64.0%;
87.5%;
                                                                                                                                                                                                                                                                   ... Dase pairs
... nucleic acid
... STRANDEDNESS: single
... TOPOLOGY: linear
US-09-237-510-6
                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 64.0
Best Local Similarity 87.5
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 62.0
Best Local Similarity 92.9
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       41 TACTCAGATGAGTCTT 26
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US-08-117-952-668/c
                                                                                                           FILING DATE:
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Patent No. 5958700

GENERAL INFORMATION:
APPLICANT: Nadeau, James G.
APPLICANT: Linn, Carl P.
APPLICANT: Linn, Carl P.
APPLICANT: Linn, Carl P.
TITLE OF INVENTION: FLUORESCENCE QUENCHING
TITLE OF INVENTION: FLUORESCENCE QUENCHING
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
ADDRESSEE: R. J. Rodrick, Becton Dickinson and Company
STREET: 1 Becton Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 12.8; DB 2; Length 48;
Pred. No. 2.7e+02;
                                                                                                                                                                                                                                                                                                                  ADDRESSEE: R. J. Rodrick, Becton Dickinson and Company STREET: 1 Becton Drive CITY: Franklin Lakes STATE: NJ COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2; Indels
                                                                                                                         GENERAL INFORMATION:
APPLICANT: Nadeau, James G.
APPLICANT: Pitner, James B.
APPLICANT: Linn, Carl P.
APPLICANT: Linn, Carl P.
TITLE OF INVENTION: DETECTION OF NUCLEIC ACIDS BY TITLE OF INVENTION: FLUORESCENCE QUENCHING NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/865,675
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                   Sequence 6, Application US/08865675 Patent No. 5928869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Fugit, Donna R.
REGISTRATION NUMBER: 32,135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 64.0%;
87.5%;
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER:
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 87.5
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 48 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3 tacccaggtgagtctt 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      41 TACTCAGATGAGTCTT 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 1 Becc...
CITY: Franklin Lakes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                           07417
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                                                             US-08-865-675-6/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-237-510-6/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-865-675-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
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US-09-101-886B-71

Sequence 71, Application US/09101886B

Patent No. 6197507

GENERAL INFORMATION:
APPLICANT: BERG, THOMAS

APPLICANT: TOLLERSRUD, OLE K

APPLICANT: NILSSEN, OIVIND

TITLE OF INVENTION: GENETIC TEST FOR ALPHA-MANNOSIDOSIS

NUMBER OF SEQUENCES: 104

CORRESPONDENCE ADDRESS:
ADDRESSEE: BARBARA G. ERNST

STREET: S55 13TH STREET, NW SUITE 701E

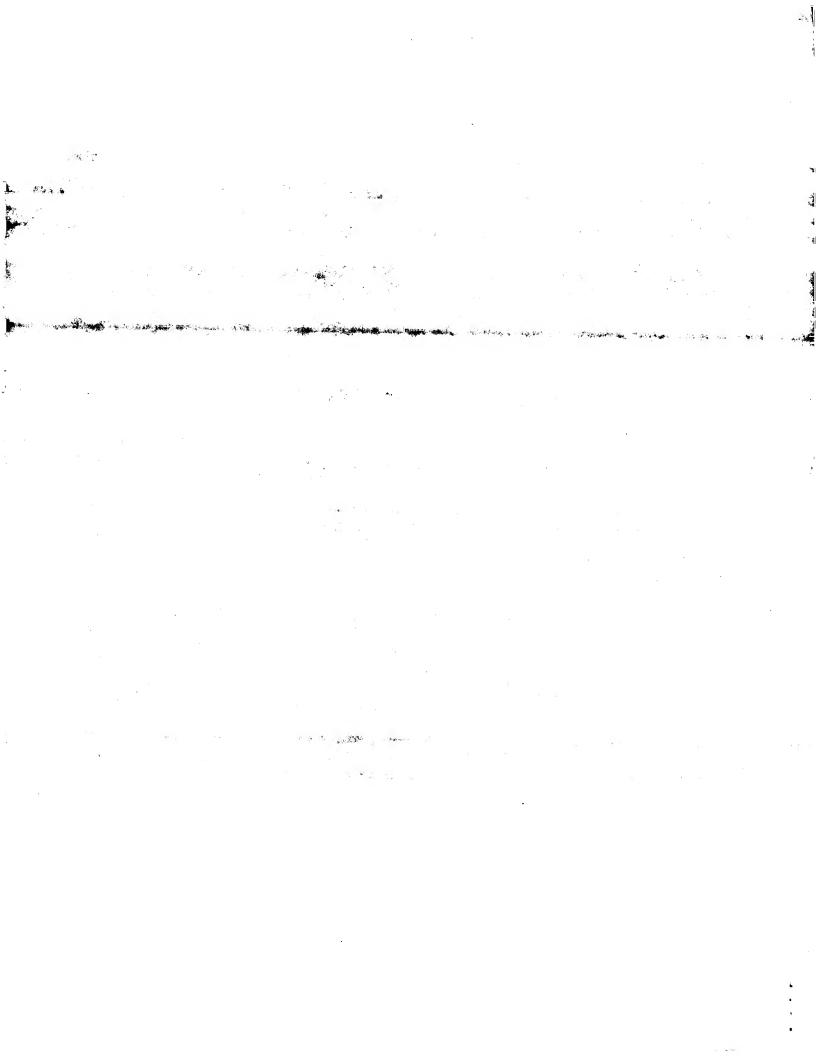
CITY: WASHINGTON

STATE: DC.
                                                                                                                                                                                                                                                                                                                                           Score 12.4; DB 4; Length 46;
Pred. No. 4.4e+02;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: RIOPPY disk
COMPUTER: IRDAPY disk
COMPUTER: IRDA PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/101,886B
FILING DATE: 29-JANUARY-1998
CLIASSIFICATION: 435
PRIOR APPLICATION NUMBER: PCT/GB97/00109
FILING DATE: 12-JANUARY-1998
CLIASSIFICATION: A35
ATTORNEY/AGENT INFORMATION:
NAME: REFERENCE/DOCKET NUMBER: 30,377
REFERENCE/DOCKET NUMBER: 31,377
REFERENCE/DOCKET NUMBER: 1181-240
TELECOMMUTICATION INFORMATION:
TELECOMMUTICATION INFORMATION:
TELECOMMUTICATION INFORMATION:
TELECOMMUTICATION INFORMATION:
TELECOMMUTICATION INFORMATION:
                                                                  860098.421
                      NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
                                                            REFERENCE/DOCKET NUMBER: 86
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
TELEPAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA (genomic)
                                                                                                                                                                                                                                                                                                                                           62.0%;
92.9%;
    ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                           LENGTH: 46 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 202-783-6031 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1: 20 base pairs
nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14 AGTACCCAAGTGAG 1
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Best Local Similarity
Matches 13; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: DAHYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                           ; TOPOLOGY:
US-09-060-410-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 12, Application US/09060410
Fatent No. 6165461
GENERAL INFORMATION:
APPLICANT: Cobb, Melanie
APPLICANT: Hutchinson, Michele
APPLICANT: Chen, Zhu
APPLICANT: Chen, Zhu
TITLE OF INVENTION: TAO PROTEIN KINASES AND METHODS OF USE
TITLE OF INVENTION: THEREFOR
TITLE OF INVENTION: THEREFOR
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
CITY: Seattle
STATE Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 12.4; DB 2; Length 22;
Pred. No. 4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                           COMPUTER FRADABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/117,952
FILING DATE: 07-SEP-1993
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/17,952
FILING DATE: 15-JUN-1993
ATTORNEY/AGENT INFORMATION:
NAME: Reiter, Stephen E.
REGISTRATION NUMBER: 31,192
REGISTRATION NUMBER: 941 9423
TELECOMMUTICATION INFORMATION:
TELECOMMUTICATION INFORMATION:
TELECOMMUTICATION OF SEQ ID NO: 668:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
FILING DATE: US/09/060,410
FILING DATE: 14-APR-1998
STREET: 444 South Flower Street, Suite 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: Oligonucleotide HYPOTHETICAL: NO
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92.9%;
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Best Local Similarity 92.9
Matches 13; Conservative
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TYPE: nucleic acid
STRANDEDNESS: single
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                      Los Angeles
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HYPOTHETICAL: NC
                                                                                  90071
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US-08-117-952-668
                                                              COUNTRY:
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Score 12.2; DB 4; Length 20;
Pred. No. 5.1e+02;
0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                             Sequence 14, Application US/08685871
Patent No. 6033499
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: RHO TARGET PROTEIN KINASE P160
NUMBER OF SEQUENCES: 68
CORRESPONDENCE ADDRESS:
ADDRESSEE: Foley & Lardner
STREET: 3000 K Street, N.W., Suite 500
CITY: Washington
STRYEE: D.C.
COUNTRY: USA
COUNTRY: USA
IP: 20007-5109
COMPUTER READABLE FORM:
MEDIUM TYPE: BM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPATIBL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUBER: US/08/685,871 FILING DATE: 24-JUL-1996 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 12.2; DB 3;
Pred. No. 5.4e+02;
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APPLICATION DATA:
APPLICATION NUMBER: JP 8-184102
FILING DATE: 25-JUN-1996
PRIOR APPLICATION DATE: 37-JUN-1996
APPLICATION NUMBER: JP 7-262553
FILING DATE: 14-SEP-1995
ATTORNEY/AGENT INFORMATION:
NAME: BENT, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DROCKET NUMBER: 16887/845
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECHANE: (202)672-5399
TELEFAX: (202)672-5399
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TELEX: 904136
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 32 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
   61.0%;
82.4%;
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82.4%;
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APPLICANT: NARUMIYA, Shuh
APPLICANT: IWAMATSU, AKIhi
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                                  Best Local Similarity 82.4
Matches 14; Conservative
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Best Local Similarity 82.4
Matches 14; Conservative
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US-08-685-871-14
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US-08-685-871-27
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                                                                                                                                                                                                                                                               OPERATION SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/685,871
FILING DATE: 24-JUL-1996
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 8-184102
FILING DATE: 25-JUN-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 7-262553
FILING DATE: 14-SEP-1995
ATTON REVIGENT INFORMATION:
NAME: BENT: Stephen A.
REGISTRATION NUMBER: 29,768
REGISTRATION NUMBER: 29,768
REGISTRATION NUMBER: 29,768
REGISTRATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFRAX: (202)672-5300
IIILE OF INVENTION: RHO TARGET PROTEIN KINASE P160
NUMBER OF SEQUENCES: 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Foley & Lardner
STREET: 3000 K Street, N.W., Suite 500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Search completed: October 2, 2001, 16:03:49
Job time: 14593 sec
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                                                                                                                                                   COUNTY: USA
21P: 20007-5109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     82.48;
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TELEX: 904136
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity 82.4
Matches 14; Conservative
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LENGTH: 32 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                           Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
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Mouse PEPCK-cytoso Chromosome 11 (loc GC PCR primer #2. GC PCR primer #2. GC PCR primer #2. GC PCR primer Strail Human retina EST 7 3' YH primer VHTP3 Physiologically ac Physiological Phy

c 12 12.4 62.0 20 22 AAF62963 Mouse PEPCK-cytose c 13 12.4 62.0 22 16 AAQ82668 Chromosome 11 (lo 14 12.4 62.0 33 16 AAQ86221 GC PCR primer #2. 15 12.4 62.0 33 16 AAQ86223 GC PCR (R) primer #2. 15 12.4 62.0 33 16 AAQ85870 Glucocerebrosidas 17 12.4 62.0 33 24 AAF98212 C neoformans stra	12.4 62.0 46 20 AAZ32444 Human retina EST 12.2 61.0 32 18 AAT78245 Physiologically 12.2 61.0 32 18 AAT78245 Physiologically 12.2 61.0 32 18 AAT78247 Physiologically	12.2 61.0 32 18 AAT/8231 Physiologically 12.2 61.0 32 18 AAT/8217 Physiologically 12.2 61.0 32 18 AAT/8217 Physiologically 12.2 61.0 32 18 AAT/8219 Physiologically 12.2 61.0 32 18.2 61	12.2 01.0 32 18 AA178221 12.2 61.0 32 18 AA178213 12.2 61.0 32 19 AAV43407 12.2 61.0 44 13 AA030902	30 12.2 61.0 44 13 31 12 60.0 20 17 32 12 60.0 20 17	12 60.0 20 19 AAV28523 12 60.0 20 20 AAV63604 12 60.0 21 20 AAX35657	36 12 60.0 21 21 AAA75059 37 12 60.0 22 22 AAA91105 38 12 60.0 23 21 AAA75062 39 12 60.0 23 21 AAA7783	12 60.0 27 12 60.0 30 12 60.0 30 12 60.0 32 11.8 59.0 17	ALIGNMENTS	RESULT 1 AAC65543 ID AAC65543 standard; DNA; 20 BP.	AC AAC65543; XX DT 12-FEB-2001 (first entry)	XX DE Human focal adhesion kinase antisense sequence #9. XX	KW Human; focal adhesion kinase; FAK; signal transduction; cancer; KW embryonic development disorder; anglogenic disorder; wound healing; KW antisense; phosphorothioate; ss.	AA Omo sapiens. XX XX PN US6133031-A.	17-OCT-2000.	PF 19-AUG-1999; 99US-0377310. XX DP 19-AUG-1999. 99US-0377310.	Hd SISI (-SISI)	AA PAI Monia BP, Gaarde WA;	DR WPI; 2001-006141/01.	PT New antisense compounds for inhibiting focal adhesion kinase PT expression, especially useful for inhibiting retinal PT neovascularization, or for diagnosing and treating e.g. colon cancer -
GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd. OM nucleic - nucleic search, using sw model	Run on: October 2, 2001, 16:18:39; Search time 876.95 Seconds (without alignments) 14.320 Million cell updates/sec	Title: US-09-757-100B-11 Perfect score: 20 Sequence: 1 agtacccaggtgagtcttag 20	Scoring table: IDENTITY_NUC Gapop 10.0 , Gapext 1.0	Searched: 730101 seqs, 313950809 residues Total number of hits satisfying chosen parameters: 854978	Minimum DB seq length: 0 Maximum DB seq length: 50	Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries	ပျ	 5: /SIDS8/gcqdata/geneseq/geneseqr/NA1984.DAT.* 6: /SIDS8/gcqdata/geneseq/geneseqn/NA1985.DAT.* 7: /SIDS8/gcqdata/geneseq/geneseqn/NA1986.DAT.* 8: /SIDS8/gcqdata/geneseqr/geneseqn/NA1987.* 	9: /SIDSB/gcgdata/geneseq/geneseqn/NA198B.DAT:* 10: /SIDSB/gcgdata/geneseq/geneseqn/NA198B.DAT:* 11: /SIDSB/gcgdata/geneseq/geneseqn/NA199D.DAT:* 12: /SIDSB/gcgdata/geneseq/geneseqn/NA199D.DAT:*	14: /SIDS8/gcgdata/geneseq/geneseqn/NA1993.DAT:* 15: /SIDS8/gcgdata/geneseq/geneseqn/NA1994.DAT:* 16: /SIDS8/gcgdata/geneseq/geneseqn/NA1994.DAT:* 16: /SIDS8/gcgdata/geneseq/geneseqn/NA1995.DAT:*		/SIDSB/gcgdata/geneseq/geneseqn/Na1999 /SIDSB/gcgdata/geneseq/geneseqn/Na2000 /SIDSB/gcgdata/geneseq/geneseqn/Na2001	Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.		Result No. Score Match Length DB ID Description	1 20 100;0 20 22 AAC65543 Humai Humai 2 75;0 15 22 AAC65563 Humai 2 75;0 21 18 AAMEG077 Humai	13.4 67.0 21 19 AAV38613 13.4 66.0 37 22 AAF92420	6 12.8 64.0 23 21 AAA72481	.0 29 19 AAV00357 Bacilor .0 29 22 AAF73275 Oligo .0 34 19 AAV15506 Prim .0 48 20 AAV82665 Targ

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human intercellular adhesion molecule-1; ICAM-1; inflammation; adult respiratory distress syndrome; multiple organ failure; GM1594;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Anti-sense oligo:nucleotide(s) for blocking ICAM-1 mRNA translation - for treating septic shock, adult respiratory distress syndrome
                                                                                                                                                                                              Antisense; pre-mRNA; mature mRNA; vascular defect; tissue defect;
                                                                                                                                                                      ICAM-1 antisense oligonucleotide #7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Column 21; 16pp; English
                                                                              1077/c
AAT58077 standard; DNA; 21
Bradley MO, Hoke GD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1997-033603/03
                                                                                                                                                                                                                                   septic shock; ss
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                                                                                                                                            18-MAR-1997
                                                                                                                                                                                                                                                                                                                                        24-JUL-1992;
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                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                  AAT58077;
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                        The present invention describes a number of phosphorothioate antisense sequences to the human focal adhesion kinase (FAK) protein. This protein is involved in integrin-mediated signal transduction, and is implicated in cancer, particularly colon, breast and oral tumours, embryonic development disorders, anglogenic disorders and wound healing. The antisense sequences, including the one shown here, can be used in the treatment of all of these.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes a number of phosphorothioate antisense sequences to the human focal adhesion kinase (FAK) protein. This protein is involved in integrin-mediated signal transduction, and is implicated in cancer, particularly colon, breast and oral tumours, embryonic autisense development disorders, angiogenic disorders and wound healing. The antisense sequences, including the one shown here, can be used in the treatment of all of these.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New antisense compounds for inhibiting focal adhesion kinase expression, especially useful for inhibiting retinal neovascularization, or for diagnosing and treating e.g. colon cancer
                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                 Human; focal adhesion kinase; FAK; signal transduction; cancer; embryonic development disorder; angiogenic disorder; wound healing; antisense; phosphorothioate; ss.
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                                                                                                                                                               DB 22; Length 20; 0.19;
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                                                                                                                             Sequence 20 BP; 5 A; 4 C; 6 G; 5 T; 0 other;
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                                                                                                                                                                                              Mismatches
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Pred. No.
 Claim 15; Column 23; 30pp; English
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                                                                                                                                                                    100.0%;
100.0%;
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Best Local Similarity 100.
Matches 20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
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Williams TJ;

Lee C,

OF NAVY.

92US-0918259. 93US-0136118. 92US-0918259.

(first entry)

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The sequences given in AAT58071-85 represent oligonucleotides which are antisense to sequences contained in the pre-mRNA or mature mRNA transcript of human intercellular adhesion molecule-1 (ICAM-1).
                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                          These oligonucleotides may be used for treating septic shock and the manifestations of septic shock, e.g. inflammation, and vascular and tissue defects. They are also useful in the treatment of septic shock associated diseases, e.g. adult respiratory distress syndrome,
                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                             Length 21;
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                                                                                                                                                                                                                                                                                             Ouery Match 67.0%; Score 13.4; DB 18; Best Local Similarity 93.3%; Pred. No. 5.3e+02; Matches 14; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                          Sequence 21 BP; 5 A; 9 C; 5 G; 2 T; 0 other;
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                                                                                                                                                                               multiple organ failure etc.
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ID AAV386
XX
AC AAV386
XX
DT 13-0CT
XX
DE Human
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Gaps

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0; Indels

75.0%; Score 15; DB 22; ilarity 100.0%; Pred. No. 76; Conservative 0; Mismatches 0;

Best Local Similarity Matches 15; Conserv

Query Match

Length 15;

gene for use in

Itoh

Suzuki M,

Ohshima T,

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This invention relates to a mouse containing the human cytochrome P450 gene (from the CYP3A family). PCR primers AAF93356 - AAF9241 are used in camples illustrating the construction of vectors used in the production of the transgenic mouse of the invention. The mouse can be used for studying human drug efficacy, metabolism and toxicity, including
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nucleotide encoding an APJ protein and/or a nucleotide encoding the CD4 protein, such that the cells coexpress APJ and CD4. The APJ receptor (AAM87466) is an orphan seven transmembrane domain receptor that has been found to function as an efficient coreceptor for cellular infection by a number of HIV-1 and SIV strains. The entry of HIV into cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to recombinant eukaryotic cells transformed with a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New recombinant eukaryotic cells coexpressing APJ and CD4 polypeptides, useful for identifying compounds that modulate interaction between an HIV virus and an APJ receptor e.g. to develop anti-HIV drugs
                                                                                                                                                                                                                                                                                                                                                                                                                                               66.0%; Score 13.2; DB 22; Length 36; 83.3%; Pred. No. 7.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Long terminal repeat; LTR; HIV-1; APJ receptor; coreceptor, cellular infection; envelope protein; env; CD4 coexpression; drug development; PCR primer; ss.
                                                                                                                                     Mouse having completely humanized human cytochrome P450 ge studying drug efficacy, metabolism and toxicity with ease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HIV-1 LTR (long terminal repeat) PCR primer, SEQ ID NO:4.
                                                                                                                                                                                                                                                                                                                                             the application of immortalised cell and tissue cultures.
                                                                                                                                                                                                                                                                                                                                                                                 Sequence 36 BP; 8 A; 7 C; 8 G; 13 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                 Example 16; Page 67; 137pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Faulds D, Hesselgesser JE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus type 1.
                                                         Kuroiwa Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 2; Page 36; 67pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99WO-EP06553
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA10299 standard; DNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 gtacccaggtgagtctta 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best_Local Similarity 83.3
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6 gtacccaagtcaggctta
                                                         Tomizuka K,
                (KIRI ) KIRIN BEER KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-256972/22.
                                                                                             WPI; 2001-202806/20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200014220-A1.
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                                                         Ishida I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA10299;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  - useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     substantially complementary to at least a portion of the pre-
or mature RNA transcript of human intracellular adhesion molecule
(ICAM). E-selectin or vascular cell adhesion molecule (VCAM).
It can be used to inhibit expression of these proteins. Inhibition
of these proteins forms the basis for treatment of conditions and
diseases that have an inflammatory component, e.g. acne, psoriasis,
arthritis, organ rejection, wounds, burns, septic shock or
inflammatory complications of septic shock.
                               vascular cell adhesion molecule-1; antisense; inflammatory; disease; treatment; septic shock; psoriasis; wounds; burns; acne; arthritis; organ rejection; inhibition; expression; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The sequence is that of an antisense oligonucleotide which is
                ICAM-1; intracellular adhesion molecule-; E-selectin; VCAM-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pro alpha 2 (I) collagen (COLIA2) specific PCR primer SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antisense oligonucleotides to ICAM-1, E-selectin or VCAM-1 for treating diseases having an inflammatory component, e.g. psoriasis, wounds and septic shock
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cytochrome P450; Cyp3A; PCR primer; transgenic mouse;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     67.0%; Score 13.4; DB 19;
93.3%; Pred. No. 5.3e+02;
live 0; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                                                 Williams TJ;
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                                                                                                                                                                                                                                                                                                                                                                                 Lee C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAF92420 standard; DNA; 36 BP
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                                                                                                                                                                                                                                                                                                                                         (DYAD-) DYAD PHARM CORP.
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                                                                                                                                                                                                                                                                                                                                                                                 Hoke GD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immortalised cell; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19 GTTCCCAGGTGAGTC
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nes 14; Conserv
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                                                                                                                                 Homo sapiens
                                                                                                                                                                           WO9824797-A1
                                                                                                                                                                                                                                                        02-DEC-1996;
                                                                                                                                                                                                                                                                                                02-DEC-1996;
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                                                                                                                                                                                                                                                                                                                                                                                 Bradley MO,
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                                                                                                                Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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Zhou Y;

Mitrovic B,

Horuk R,

ð g WPI; 2000-543395/49.

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involves binding of the viral envelope protein (env) to CD4, followed by interaction with a coreceptor. Binding to the coreceptor triggers a conformational change in env that mediates fusion between the viral membrane and the host cell membrane. The invention also encompasses peptides and antibodies which specifically bind to an extracellular domain of APJ. These inhibit membrane fusion between a cell coexpressing the APJ and CD4 proteins and a cell expressing an HIV env protein and the APJ and a cell expressing and the APJ inhibits HIV infection of the APJ/CD4-expressing cell. The new recombinant cells provide an important tool for investigating and controlling HIV infection. The cells can be used to identify compounds that modulate interaction between HIV and an APJ receptor, which may be useful in the development of anti-HIV drugs. Contacting cells expressing APJ and CD4 with an APJ binding/Dlocking agent (e.g., the antibodies or peptides) may be useful to inhibit HIV infection of such cells, e.g., to treat subjects having an HIV-related disorder associated with APJ expression. Antibodies and peptide fragments can be included in medians and administered to treat patients (especially foctuses) having, or at risk of developing, an HIV infection or related disorder. The antibodies can also be used detect cells expressing the APJ receptor, which are useful to diagnose susceptibility to HIV infection or related disorder. The art subjects are useful to diagnose susceptibility to medicion or related disorder. The art used to a susceptibility to the APJ receptor.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               higher APJ levels in central nervous system tissues may indicate an increased risk of neuropathogenesis associated with HTV infection. Transgenic animals which express human CD4 and APD proteins may provide model systems for the study of HTV infection and for anti-HTV drug development. Sequences AAA10299-A10300 represent PCR primers used in an terminal repeat) DNA sequences from CD4/APJ-expressing cells which had been previously exposed to HTV-1. The amplified LTR DNA was detected via
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Soybean; chalcone reductase; phenylpropanoid pathway; isoflavone synthase; isoflavonoid biosynthesis; defence response; attractant; repellant; signal compound; antibiotic; transgenic plant; transgenic seed; PCR primer; ss.
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Soybean chalcone reductase PCR primer, SEQ ID NO:65.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hybridisation with a labelled probe (AAA10301)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 23 BP; 7 A; 7 C; 5 G; 4 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (DUPO ) DU PONT DE NEMOURS & CO E I.
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99US-0144783.
99US-0156094.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  64.0%;
87.5%;
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Best Local Similarity 87.5
Matches 14; Conservative
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Sequences AAA72437, AAA72449-A72461, AAA72468 and AAA72474-A72477

represent cDNAs encoding novel plant isoflavone synthases (AAB21052 and AAB21054- AAB21071). AASO discolosed is the soybean cytochrome P450

monocoxygenase CYP93C1, encoded by a known sequence (AAA72444, NCBI No. 2739005), which was identified in the present invention as having isoflavone synthase expression constructs, transformed host cells, and transgenic plants and seeds conscribing the novel CDNA sequences of the invention. The invention also constructs, transformed host cells, and transgenic plants and seeds conscribing isoflavone synthase expression in a host cell, altering isoflavoned in a plant, and identifying nucleic acids encoding other plant isoflavone synthases. Isoflavone synthase cells, altering isoflavonoid levels in a plant, and identifying nucleic acids encoding other plant isoflavone synthases. Isoflavonoids are cells as key roll in the biosynthesis of isoflavonoids is soflavonoids are a class of secondary metabolites mainly produced in leguminous plants by a branch of the phenylpropanoid pathway. Isoflavone synthase catalyses the first step in the branch of this pathway that commits metabolic intermediates to the synthesis of isoflavonoids. Isoflavonoids is participate in the defence response of legumes against phytopathogenic intermediates to the synthesis of isoflavonoids. Isoflavonoids consumption of legumes and rhizobial bacteria which eventually result in nodulation and nitrogen fixation. They have also been shown to act as artibiotics, repellents, attractants, and signal compounds, and consumption of legume isoflavonoid synthases, and nucleic acids encoding them are useful for altering the levels of isoflavonoids produced in legumes such as soybean, and for the production of isoflavonoids produced in altering the levels of isoflavonoids produced in legumes such as soybean, and for the production of isoflavonoids camplification of the invention to amplify DNA encoding synthase; in the actual actual actual actual actual 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                            Nucleic acids encoding isoflavonoid synthases, useful for producing transgenic plants with increased production of isoflavonoids which are involved in defense against phytopathogenic microorganisms -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Insecticidal protein; Bacillus thuringiensis; monocotyledonous plant; structural gene; maize; CryI(b); CryIIB; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bacillus thuringiensis insecticidal gene modification primer BTK58.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 25 BP; 6 A; 7 C; 5 G; 7 T; 0 other;
                                                                                                                                          Example 14; Page 52; 157pp; English.
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Bacillus thuringiensis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
Matches 14; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JS5689052-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-NOV-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAV00357;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAV00357/c
ID AAV003
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Gaps

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Length 29; Indels

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The present sequence was used in the development of a novel system, comprising an adeno-associated virus (AAV) vector containing a foreign DNA, and the rep 68/78 sequence of AAV, the expression of which is delayed. The components may be present in cis (in a single agent) or in trans (in separate agents). The system is used for production of AAV vectors, particularly for gene therapy. The foreign DNA may encode a therapeutic protein, e.g. interferon, interleukin, growth factor, coagulation factor or metabolic enzyme, particularly one that increases the immunogenicity of tumour cells, and/or a diagnostic protein. The system produces AAV vectors on a large scale. It is based on the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 discovery that the rep68 and 78 proteins interfere with replication of AAV DNA, and that this interference is overcome by delaying expression of these proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   System of adeno-associated viral vector and rep 68/78 sequence of this virus - expression of which is delayed until replication of viral DNA has started, provides large scale production of vectors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 64.0%; Score 12.8; DB 19; Length 34; Best Local Similarity 87.5%; Pred. No. 1.1e+03; Matches 14; Conservative 0; Mismatches 2; Indels C
                                                                                                                                                                                                                                                                                                                                                                         AAV; vector; rep 68/78; PCR primer; gene therapy; ss.
                                                                             Score 12.8; DB 22;
Pred. No. 1.1e+03;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 34 BP; 4 A; 13 C; 9 G; 8 T; 0 other;
                                Sequence 29 BP; 6 A; 9 C; 6 G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                            Primer for adeno-associated virus DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Maass G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example; Page 7; 20pp; German.
                                                                                                                                                                                                                                                AAV15506 standard; DNA; 34 BP
                                                                              64.0%;
87.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 97WO-DE01333.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             96DE-1025188
 Bacillus thuringiensis.
                                                                                                                                                                                                                                                                                                               (first entry)
                                                                             Query Match 64.0
Best Local Similarity 87.5
Matches 14; Conservative
                                                                                                                                          1 agtacccaggtgagtc 16
                                                                                                                                                                      6
                                                                                                                                                                                                                                                                                                                                                                                                                      Adeno associated virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bogedain C, Hallek M,
                                                                                                                                                           24 AGTGCCCAGGAGAGTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (MEDI-) MEDIGENE GMBH. (MEDI-) MEDIGENE AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1998-086641/08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    for gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9749824-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-JUN-1996;
                                                                                                                                                                                                                                                                                                             11-JUN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31-DEC-1997
                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                AAV15506;
                                                                                                                                                                                                                                  AAV15506
                                                                                                                                                                                                                     RESULT
   SXC
                                                                                                                                           δλ
                                                                                                                                                                      윤
                                                                                                                                                                                                                                                                               ó
                                                                                                                                                                                                               The present sequence represents a primer used in the present invention describing new structural genes capable of being expressed in a monocotyledonous plant. The new genes comprise modified nucleotide sequences which encode insecticidal proteins of Bacillus thuringiensis. The genes have been modified to reduce the usage of codons that are rare or semi-rare in monocotyledon DNA, thereby increasing transformation efficiency and/or increasing accumulation of the insecticidal protein in monocotyledon tissues.
                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel nucleic acids, useful for transgenic plant production which is capable of expressing increased levels of desired proteins -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to nucleotides 669-1348 of a B.thuringiensis CryIA(b). The invention is useful for transgenic plant production, e.g. maize, capable of expressing increased amount of transgenic protein, e.g. crystal protein toxin gene
                                                                                                                                                                                                                                                                                                                                                                                                                                   ;
0
                                                                                                                                        encoding insecticidal proteins of Bacillus thuringiensis led to enhance expression in monocotyledonous plants
                                                                                                                                                                                                                                                                                                                                                                                                      Length 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CryIA; transgenic; crystal; toxin; insecticide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                    Score 12.8; DB 19;
Pred. No. 1.1e+03;
); Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                       Sequence 29 BP; 6 A; 9 C; 6 G; 8 T; 0 other;
                                                                             Sanders PR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sanders PR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Column 16; 81pp; English.
                                                                                                                                                                                   Example 1; Column 17; 86pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                   :
                                                                           Brown SM, Dean DA, Fromm ME,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Fromm ME,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
                                                                                                                                                                                                                                                                                                                                                                                                  Ouery Match
Best Local Similarity 87.5%;
Matches 14; Conservative
 93US-0172333
                95US-0530492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         95US-0530492
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAF73275 standard; DNA; 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Brown SM, Dean DA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-190861/19.
                                            (MONS ) MONSANTO CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MONS ) MONSANTO CO.
                                                                                                        WPI; 1998-008070/01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligonucleotide #66
22-DEC-1993;
            19-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US6180774-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-SEP-1995;
22-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                        Genes enc
modified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAF73275;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF73275/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT
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Gaps

ð g AAV82665;

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The present sequence is one of a number of antisense compounds of up to phosphoenol pyruvate carboxykinase-cytosolic (PEPCK-cytosolic). The antisense compounds are useful for inhibiting the expression of PEPCK-cytosolic in cells or tissues. They are commonly used as research reagents and in diagnostics, e.g. to elucidate the function of particular genes. They are also useful for distinguishing between functions of various members of a biological pathway and for research use. The antisense compounds are also useful prophylactically, e.g. to prevent or delay infection, inflammation or tumour formation. The present sequence is a chimeric phosphorothioate oligonucleotide with 2'-MOE wings and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antisense compound capable of modulating the expression of phosphoenol pyruvate carboxykinase-cytosolic, useful for preventing or delaying infection, inflammation or tumor formation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence sampled mapping; genomic analysis; complex genome mapping; cosmid library; chromosome 11; sequence tagged site; STS analysis; ss.
                                                                                                                                                    Mouse; antiinflammatory; cytostatic; antisense gene therapy; phosphoenol pyruvate carboxykinase-cytosolic; PEPCK-cytosolic; infection; inflammation; tumour formation; phosphorothioate; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 62.0%; Score 12.4; DB 22; Length 20; Ilarity 92.9%; Pred. No. 1.8e+03; Conservative 0; Mismatches 1; Indels (
                                                                                                              Mouse PEPCK-cytosolic antisense oligonucleotide ISIS 113360.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chromosome 11 (locus LDHC) STS primer LDHC-Z.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 5 A; 6 C; 5 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 17; Column 44; 64pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Wyatt J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
                                                                                                                                                                                                                                                                                                                                                                          21-JAN-2000; 2000US-0488671.
                                                                                                                                                                                                                                                                                                                                                                                                                   21-JAN-2000; 2000US-0488671.
                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ82668 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cccaggtgagtctt 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Butler MM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-190979/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15 CCAAGGTGAGTCTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
Matches 13; Conserv
                                                                                                                                                                                                                                                                                       US6187545-B1
                                                                   08-MAY-2001
                                                                                                                                                                                                                                                 Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-SEP-1995
                                                                                                                                                                                                                                                                                                                                  13-FEB-2001
                           AAF62963;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ82668;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        McKay R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ß
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ82668/
      ογ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                detector oligonucleotide of the invention. The detector oligonucleotide comprises a single-stranded target-binding region and an intramolecularly base-paired secondary structure linked to two dyes (donor and acceptor fluorophores). In the secondary structure, fluorescence of the donor is quenched, but when it is linearised or unfolded a change in some fluorescence parameter becomes detectable. A target nucleic acid is detected by hybridizing it to a detectable. A oligonucleotide in which the secondary structure is 5' to the target binding region, primer extension to produce a complementary strand using the secondary structure as template, resulting in linearization or oligonucleotide are used to detect (by primer extension and hybridization) and amplify (as signal primer) target sequences, e.g. for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Target oligonucleotides AAV82665-68 were designed to hybridise to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                               Detector oligonucleotide; hairpin structure; fluorescence; primer extension; hybridization; signal primer; frameshift mutation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New detector oligo:nucleotide having base-paired region carrying quenched dyes - where dyes become fluorescent when region is unpaired, useful for detection and amplification of target nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 12.8; DB 20;
Pred. No. 1.2e+03;
); Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 48 BP; 15 A; 8 C; 10 G; 15 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Schram JL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2; Page 12; 20pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nadeau JG, Pitner BJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     detecting frameshift mutations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 12
AAF62963/c
ID AAF62963 standard; DNA; 20 BP.
                                                                                                                                                           BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (BECT ) BECTON DICKINSON & CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cch 64.0%;
il Similarity 87.5%;
14; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98EP-0109682
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          97US-0865675
                                                                                                                                       AAV82665/c
ID AAV82665 standard; DNA; 48
                                                                                                                                                                                                                                                                                       Target oligonucleotide JN5
                                                                                                                                                                                                                                            (first entry)
3 tacccaggtgagtct 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           41 TACTCAGATGAGTCTT 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1999-001406/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-MAY-1997;
                                                                                                                                                                                                                                            25-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                              EP881302-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-DEC-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
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Linn PC,

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Gaps

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Synthetic

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15-JUN-1994; 15-JUN-1993; 07-SEP-1993;

WO9429486-A.

22-DEC-1994

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The sequences given in AAQ86220-21 are primers which were used in the construction of a Sindbis glucocerebrosidase (GC) vector producing cell line. These primers amplify a 521 bp fragment for GC screening of the transformant cell lines. The cell lines contain alphavirus vectors which express multiple heterologous genes. The vectors are eukaryotic layered vector initiation systems (ELVIS) derived from Sindbis. ELVIS's comprise a 5' sequence capable of initiating transcription of an alphavirus, a nucleotide sequence encoding alphavirus non-structural proteins, a viral junction region which has been inactivated such that viral transcription of the subgenomic fragment is prevented, and an alphavirus RNA polymerase recognition sequence. Inactivation of the viral junction region prevents transcription of the subgenomic fragment making vectors such as this transcription of systic fibrosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryotic layered vector initiation system; Sindbis; alphavirus; PCR; transcription initiation; non-structural protein; subgenomic fragment; RNA polymerase recognition sequence; ELVIS; gene therapy; amplify; primer; polymerase obinin reaction; cystic fibrosis; glucocerebrosidase; ss.
                                                                                                                                                                                                               New alpha virus vectors for gene therapy - of viral infection, cancer, auto:immune disease, etc., and as vaccines.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 12.4; DB 16; Length 33;
Pred. No. 1.8e+03;
0; Mismatches 1; Indels
                                                                                                           Dubensky TW, Ibanez CE, Jolly DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dubensky TW, Ibanez CE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 33 BP; 10 A; 5 C; 11 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                           Example 13; Page 154; 260pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ86223 standard; DNA; 33 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     62.0%;
92.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94WO-US10469.
93US-0122791.
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94US-0198450
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-DEC-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 92.9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chang SMW, Driver DA,
                                                                                                      Chang SMW, Driver DA,
Polo JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6 agtagccaggtgag 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 agtacccaggtgag 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GC PCR (R) primer #2.
                                                                                                                                                                       WPI; 1995-131362/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (VIAG-) VIAGENE INC.
                                                              (VIAG-) VIAGENE INC
15-SEP-1993;
18-FEB-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-SEP-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-FEB-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cosmids by automated sequencing without intermediate subcloning.

A sample of 371 DNA sequence fragments were determined and of these, 277 were suitable for STS primer prediction by computer analysis (using the "Primer" program available from E.Lander, MIT).

The STSs and cosmids were mapped by in situ hybridisation, somatic cell hybrid analysis or both. Using this method, 370 STSs specific for human chromosome 11 were generated and most of them were regionally mapped; This procedure illustrates a novel method for sequencing complex genomes, designated "sequence sampled mapping".

The sequence sampled mapping method is useful for the completion of high density sequence-based maps, and ultimately, for the complete sequencing of genomic DNA directly from cosmid clones.

See AAQ82001-Q82706 and AAQ91325-Q91358 for STS primers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryotic layered vector initiation system; Sindbis; alphavirus; PCR; transcription initiation; non-structural protein; subgenomic fragment; RNA polymerase recognition sequence; ELVIS; gene therapy; amplify; primer; polymerase obain reaction; cystic fibrosis; glucocerebrosidase; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequences were determined from the ends of chromosome 11-specific
                                                                                                                                                                                                                                                                                                                    Sequencing complex genomes, present as fragments in a cosmid
library - by sequencing end-specific nucleotides of each clone
then correlating with spatial relationship of cosmid, esp. for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 12.4; DB 16;
Pred. No. 1.8e+03;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 22 BP; 8 A; 5 C; 4 G; 5 T; 0 other;
                                                                                                                                                                                       (SALK ) SALK INST BIOLOGICAL STUDIES.
                                                                                                                                                                                                                                                                                                                                                                                                                          Example 4; Page 92; 128pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ86221 standard; DNA; 33 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  62.0%;
92.9%;
                                                                                                                        93US-0078471.
93US-0117952.
                                                                                  94WO-US06810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94WO-US10469.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-DEC-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 92.9
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mammalian chromosomes
                                                                                                                                                                                                                                   Evans GA, Smith MW;
                                                                                                                                                                                                                                                                           WPI; 1995-036508/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GC PCR primer #2.
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Gaps ö

Jolly DJ;

15-SEP-1994;

WO9507994-A

Synthetic

AAQ86221;

RESULT 14 AAQ86221

ö q 23-MAR-1995

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PI Polo JM;

XX

XX

XX

XX

New alpha virus vectors for gene therapy - of viral infection,

PT cancer, auto:immune disease, etc., and as vaccines.

XX

XX

Example 17; Page 166; 260pp; English.

XX

CC construction of a Sindbis glucocerebrosidase (GC) vector producing

CC construction of a Sindbis glucocerebrosidase (GC) vector producing

CC cell line. These primers amplify a 521 bp fragment for GC screening

CC the transformant cell lines. The cell lines contain alphavirus

CC cell line. These primers amplify a 521 bp fragment for GC screening

CC the transformant cell lines. The cell lines contain alphavirus

CC eukaryotic layered vector initiation systems (ELVIS) derived from

CC sindbis. ELVIS's comprise a 5's equence capable of initiating

CC transcription of an alphavirus, a nucleotide sequence encoding

CC alphavirus non-structural proteins, a viral junction region winch

CC has been inactivated such that viral transcription of the subgenomic

CC fragment is prevented, and an alphavirus RNA polymerase recognition

CC sequence. Inactivation of the viral junction region prevents

CC transcription of the subgenomic fragment making vectors such as this

CC suitable for a wide variety of applications, eg. gene therapy for the

XX

Sequence 33 BP; 10 A; 5 C; 11 G; 7 T; 0 other;
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Query Match 62.0%; Score 12.4; DB 16; Length 33; Best Local Similarity 92.9%; Pred. No. 1.8e+03; Matches 13; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Db 6 agtagccaggtgag 19

Search completed: October 2, 2001, 16:18:40 Job time: 15484 sec

Appli Appli Appli

Sequence Sequence Sequence

Appl Appl Appl Appl Appl Appli Appli

Sequence Sequence Sequence S

Sequence

Sequence

Sequence Seq

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Sednence:

Searched:

Database

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APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310B
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 10
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310B
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver: 2.0
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ó,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 20; DB 3;
100.0%; Pred. No. 0.12;
ive 0; Mismatches 0;
US-08-394-232A-4
PCT-US95-04080-3
PCT-US95-04080-3
US-07-603-451A-10
US-08-606-822A-11
US-08-606-822A-12
US-08-606-822A-19
PCT-US94-05257-19
US-08-606-822A-19
PCT-US94-05257-19
US-08-844-655-15
US-08-8470-187-9
US-08-318-905-25
US-08-4170-187-9
US-08-418-15-28
US-08-418-15-28
US-08-418-15-28
US-08-483-140-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                , OTHER INFORMATION: antisense sequence US-09-377-310-10
                                                                                                                                                                                                                                                                                                                        ; Sequence 10, Application US/09377310B; Patent No. 6133031; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; Sequence 30, Application US/09377310B
; Patent No. 6133031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 100.0%; P.
Watches 20; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 ggtaagcagctgccattatt 20
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    4444444444
  US-09-377-310-10
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  TYPE: DNA
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    000000000
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Appli
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Sequence 185, App
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Sequence 13, Appl
Sequence 28, Appl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Appl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 9, Appli
Sequence 5, Appli
                                                                                      Search time 417.38 Seconds (without alignments) 9.071 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 45,
Sequence 45,
Sequence 6, A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 3, P
Sequence 4, P
Sequence 3, P
Sequence 4, P
Sequence 3, P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 6,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Description
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                                                                                                                                                                                                                                                                                                                                                                                        /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
            GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-471-371-6
US-08-833-37-5
US-08-833-37-5
US-08-185-286-45
US-09-106-216-52
US-09-106-216-53
US-09-106-216-53
US-09-106-216-53
US-09-106-216-53
US-09-106-216-53
US-08-12-929-22
US-08-72-929-22
US-08-772-929-22
US-08-772-920-5
US-08-617-454-9
US-08-617-454-9
US-08-617-454-9
US-09-157-206-5
US-09-417-863-5
                                                                                                                                                                                                                                                         hits satisfying chosen parameters:
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US-08-394-232A-3
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US-09-135-642-3
                                                                                                                                                                                                                                  324599 segs, 94655562 residues
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                                                                                       2, 2001, 16:03:48
                                                                                                                                                                                                                                                                                                                        Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                               - nucleic search, using sw model
                                                                                                                                                                    1 ggtaagcagctgccattatt 20
                                                                                                                                                                                            IDENTITY_NUC
Gapop 10.0 , Gapext 1.0
                                                                                                                                                                                                                                                                                                                                                                              Issued_Patents_NA:*
                                                                                                                                     US-09-757-100B-10
20
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Match Length
                                                                                                                                                                                                                                                                                   Minimum DB seq length: 0
Maximum DB seq length: 50
                                                                                         October
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RIBOSOME-INACTIVATING PROTEINS, INACTIVE
PRECURSOR FORMS THEREOF, A PROCESS FOR MAKING A METHOD OF
USING
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/833,377 FILING DATE: 04-APR-1997 CLASSIFICATION: 435 PRIOR APPLICATION NUMBER: US 08/510,736 FILING DATE: 04-APG-1995 ATTORNEY/AGENT INFORMATION:
NAME: Stochalla, John R.
REGISTRATION NUMBER: 32,944 REFERENCE/DOCKET NUMBER: 32,944 REFERENCE/DOCKET NUMBER: 015389-001110US
                                                                                                                                                                     ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                    APPLICANT: Vasserot, Alain P.
APPLICANT: Pruzan, Ronald A.
APPLICANT: Kealey, James T.
TITLE OF INVENTION: Purified Telomerase
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "011go 13'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mod_base= OTHER
                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 45, Application US/08378761A
Patent No. 5635384
GENERAL INFORMATION:
APPLICANT: WALSH, TERENCE A
APPLICANT: HEY, TIMOTHY D
APPLICANT: MORGAN, ALICE ER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SED ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 ggtaagcagctgccattatt 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: modified_base LOCATION: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 80.03
Matches 16; Conservative
                   Lichtsteiner,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGIH: 30 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
                                                                                                                                                                                                              CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: FITTLE OF INVENTION: FITTLE OF INVENTION: UNMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: 1
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; LOCATION: 1..30
; OTHER INFORMATION:
US-08-833-377-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
                                                                                                                                                                                                                                                                         ZIP: 94111-3834
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE:
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                                                                                                                                               0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: CARELLA, BYRNE, BAIN, GILFILLAN, CECCHI, ADDRESSEE: STEWART & OLSTEIN
STREET: 6 Becker Farm Road
CITY: Roseland
STATE: N.J.
COUNTRY: U.S.A
ZIP: 07068
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
FILING DATE: 06-JUN-1995
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                          RESULT 3
US-08-471-371-6
; Sequence 6, Application US/08471371
; Patent No. 5981215
; GENERAL INFORMATION:
APPLICANT: Meissner, Paul S.
APPLICANT: Coleman, Timothy A.
TITLE OF INVENTION: HUMAN CRIPTIN GROWTH FACTOR; NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          69.0%; Score 13.8; DB 2;
88.2%; Pred. No. 1.7e+02;
iive 0; Mismatches 2;
                                                                                                      75.0%; Score 15; DB 3; 100.0%; Pred. No. 38;
                                                                                                                                               0; Mismatches
; FEATURE:
; OTHER INFORMATION: antisense sequence
US-09-377-310-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .....rybE: other nucleic acid
bESCRIPTION: /desc = "PRIMER"
US-08-471-371-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATOCKNEY/ACEMT INPORMATION:
NAME: FECTATO Gregory D.
REGISTRATION NUMBER: 36.134
REFERENCE/DOCKET NUMBER: 3258:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (201) 994-1704
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 5, Application US/08833377; Patent No. 5968506; GENERAL INFORMATION: APPLICANT: Weinrich, Scott L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 69.0
Best Local Similarity 88.2
Matches 15; Conservative
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                                                                                                                                               15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 30 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                              3 taagcagctgccatt 17
                                                                                                                         Best Local Similarity
Matches 15; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
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                                                                                                      Query Match
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Gaps

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APPLICANT: Griffin, John H
APPLICANT: Griffin, John H
APPLICANT: Bouma, Bonno N
APPLICANT: Bouma, Bonno N
APPLICANT: Bouma, Borno N
TITLE OF INVENTION: RECEIVED
TITLE OF INVENTION: IN C4DP BINDING ACTIVITY, COMPOSITIONS AND THERAPEUTIC
TITLE OF INVENTION: METHODS
TOTALE OF INVENTION: METHODS
CORRESPONDENCE ADDRESS:
ADDRESSEE: The Scripps Research Institute, Office of ADDRESSE: Patent Counsel
STREET: 10666 No. 5405946th Torrey Pines Road, TPC 8
                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                         67.0%; Score 13.4; DB 1; Length 18; 93.3%; Pred. No. 2.5e+02; live 0; Mismatches 1; Indels
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Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/985,691
FILING DATE: 19921202
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: FILLING, THOMBER: TSR0042P
REGISTRATION NUMBER: TSR0042P
TELEPHONE: 619-554-2937
TELEPHONE: 619-554-6312
INFORMATION FOR SEQ ID NO: 6:
NAME: BORUCKI, ....
REGISTRATION NUMBER: 33651
REFERENCE/DOCKET NUMBER: 3827;
TELECOMMUNICATION INFORMATION:
TELEPHONE: (317) 337-4846
INFORMATION FOR SEQ ID NO: 45:
SEGUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 6, Application US/07985691; Patent No. 5405946; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 92037
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA (genomic)
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83.3%;
                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 93.3
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 30 base pairs
TYPE: NUCLEIC ACID
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                   3 taagcagctgccatt 17
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Best Local Similarity
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MOLECULE TYPE: DN
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US-07-985-691-6
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APPLICANT: HEY, TIMOTHY D
APPLICANT: MORGAN, ALICE ER
TITLE OF INVENTION: RIBOSOME-INACTIVATING PROTEINS, INACTIVE
TITLE OF INVENTION: USING
MUMBER OF SEQUENCES: 81
CORRESPONDENCE ADDRESS:
ADDRESSE: ANDREST: ADDRESS:
CORRESPONDENCE ADDRESS:
ADDRESSE: ANDREST: STREET: 9330 ZIONSVILLE ROAD
CITY: INITANAPOLIS
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                                                                                                                                                                             MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC COMPATIBLE

COMPUTER: IBM PC COMPATIBLE

COMPUTER: IBM PC COMPATIBLE

COMPUTER: IBM PC COMPATIBLE

OFFRATION SYSTEM: PC-DOS/MS-DOS

SOFFWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/378,761A

FILING DATE: 26-JAN-1995

CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:

NAME: BORUCKI, ANDREA T

REGISCHAATION NUMBER: 33651

REFERENCE/DOCKET NUMBER: 33651

TELEPHONE: (317) 337-4846

INFORMATION FOR SEQ ID NO: 45:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 13.4; DB 1;
Pred. No. 2.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ver
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/378761
FILING DATE: 26-JAN-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 45, Application US/08485286
Patent No. 5646026
Patent No. 5646026 5646119
                                  ADDRESSEE: ANDREA T. BORUCKI
STREET: 9330 ZIONSVILLE ROAD
CITY: INDIANAPOLLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     67.0%;
93.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 67.0°
Best Local Similarity 93.3°
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                   COUNTRY: US
ZIP: 46268
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3 taagcagctgccatt 17
             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17 TAAGCAGCTGCAATT 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-485-286-45/c
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Sequence 374, Application US/09042353
Fatent No. 6255458
GENERAL INFORMATION:
APPLICANT: Lonberg, Nils
APPLICANT: Ray, Robert M.
TITLE OF INVENTION: Transgenic No. 6255458-Human Animals for TITLE OF INVENTION: Producing Heterologous Antibodies
NUMBER OF SEQUENCES: 421
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LIP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
ZIP: 94111-3834
COMPOTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPOTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/AKS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DAFA:
APPLICATION NUMBER: US/09/042,353
FILING DATE: 13-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO PCT/US96/16433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/990,860
FILING DATE: 16-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/053,131
FILING DATE: 26-APR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/096,762
FILING DATE: 22-UL-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/096,762
FILING DATE: 18-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/810,279
FILING DATE: 17-DEC-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/853,408
FILING DATE: 18-MAR-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/904,068
FILING DATE: 23-UUN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 09-WAR-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 10-OCT-1995
PRIOR APPLICATION DATA:
APPLICATION DATA:
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FILING DATE: 03-DEC-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/165,699
FILING DATE: 10-DEC-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 10-OCT-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/758,417
FILING DATE: 02-DEC-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US97/21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 08/352,322 FILING DATE: 07-DEC-1994
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     UMBER: US 08/728,463
10-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 18-NOV-1
PRIOR APPLICATION DATA:
APPLICATION NUMBER: U
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                         US-09-042-353-374/C
    RESULT
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APPLICANT: Jeunemairre, Xavier
APPLICANT: Jeunemairre, Xavier
APPLICANT: Journal Florent
APPLICANT: Lofton, Richard P.
APPLICANT: Corvol, Pierre
APPLICANT: Corvol, Pierre
APPLICANT: Corvol, Pierre
APPLICANT: Corvol, Pierre
APPLICANT: Sources, Yuri
APPLICANT: Sources, Yuri
APPLICANT: Corvol, Pierre
APPLICANT: Sources, Yuri
APPLICANT: Sources, Yuri
APPLICANT: Sources, Yuri
APPLICANT: Sources, Yuri
APPLICANT: Florent Pierre
APPLICANT: Logo, Ernst & Kurz
ADDRESSEE: Rothwell, Flgg, Ernst & Kurz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 29;
    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Rothwell, Figg, Ernst & Kurz
STREET: 555 Thirteenth Street N.W., Suite 701-E
CITY: Washington
STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC COMPATLBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 12.6; DB 3;
Pred. No. 6.6e+02;
0; Mismatches 4;
  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 09/
FILING DATE: 08-JUN-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/319,545
FILING DATE: 07-CCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/952,545
FILING DATE: 30-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Innen, Jeffrey L.
REGISTATION NUMBER: 28,937
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/106,216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2323-124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: other nucleic acid DESCRIPTION: /desc = "primer"
                                                                                                                                                                              US-09-106-216-52/c , Sequence 52, Application US/09106216 ; Patent No. 6153386 ; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ..
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-783-6040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            63.0%;
78.9%;
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                                              26 GGAAGAGCTGCCGTTGTT 8
                                                                                        20 TGAGCAGCTTCCTTTATT 3
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SEQUENCE CHARACTERISTICS:
LENGTH: 29 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 63.0
Best Local Similarity 78.9
Matches 15; Conservative
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DESCRIPTION:
US-09-106-216-52
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Juan, Shao-Chieh
APPLICANT: Lichenstein, Henri S.
APPLICANT: Lichenstein, Henri S.
APPLICANT: Lichenstein, Henri S.
TITLE OF INVENTION: ANTI-INFLAMMATORY CD14 POLYPEPTIDES
UNDMER OF SEQUENCES: 38
CORRESPONDENCES: ADGRESS:
ADDRESSEE: AMGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: US
ZIP: 91320-1789
COMPUTER READABLE FORM:
MEDIOM TYPE: FLOPPY disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 12.4; DB 2;
Pred. No. 8.7e+02;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PAPLICATION NUMBER: US/08/484,397A
FILING DATE: 07-JUN-1995
CLASSIFICATION: 124
ATTORNEY/AGENT INFORMATION:
NAME: CULTY, Daniel R.
REGISTRATION NUMBER: 32,727
REFERENCE/DOCKET NUMBER: A-324A
TELECOMMUNICATION INFORMATION:
TELEPHONE: 805/447-1000
                                                                                                                      MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "primer"
US-09-106-216-53
                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Sequence 24, Application US/08484397A
; Patent No. 5869055
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ....ORESSEE: AMGEN INC
STREET: 1840 DeHavilland Drive
CITY: Thousand Oaks
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  62.0%;
92.9%;
                                                                                                                                                                                                                                                                                                             2 gtaagcagctgccattatt 20
                                                                                                                                                                                                                                                                                                                                                 46 GGAAGAAGCTGCCGTTGTT 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 805/447-1090 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                       Query Match
Best Local Similarity 78.9°
Matches 15; Conservative
                   SEQUENCE CHARACTERISTICS:
LENGTH: 49 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 92.9
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  805/447-1090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 39 base pairs
  INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5 agcagctgccatta 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24 AGCAGCTGCCTTTA 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                       linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; MOLECULE TYPE:
US-08-484-397A-24
                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 11
US-08-484-397A-24/c
                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT
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                                                                                                                                                                                                                                                                                                                                                                         DB 4; Length 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Lalouel, Jean-Marc
APPLICANT: Jeunemaitre, Xavier
APPLICANT: Jeunemaitre, Xavier
APPLICANT: Lofton, Richard P.
APPLICANT: Soubrie, Florent
APPLICANT: Societysev, Yuri
APPLICANT: Corvol, Pierre
TITLE OF INVENTION: Method to Determine Predisposition to
TITLE OF INVENTION: Hypertension
NUMBER OF SEQUENCES: 58
                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Rothwell, Figg, Ernst & Kurz
STREET: 555 Thirteenth Street N.W., Suite 701-E
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 20004

CMDUTER READABLE FORM:
MEDIUM TYEE Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/106,216
                                                                                                                                                                                                                                                                                                                                                                    Score 12.6; DB 4
Pred. No. 7e+02;
0; Mismatches
                                    NAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 014643-009040US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08 09/
FILING DATE: 08-UNN-1998
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/319,545
FILING DATE: 07-CCT-1994
PRIOR APPLICATION NUMBER: US 07/952,545
FILING DATE: 30-SEP-1992
ATTORNEY AGENT INFORMATION:
NAME: INDEM: JEFF 1.
RECISTRATION NUMBER: 28,957
REFERENCE/DOCKET NUMBER: 2323-124
TELECHOME: 202-783-6040
TELEFAX: 202-783-6040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 53, Application US/09106216
Patent No. 6153386
GENERAL INFORMATION:
                                                                                            TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
INFORMATION FOR SEQ ID NO: 374:
SEQUENCE CHARACTERISTICS:
LENGTH: 44 base pairs
                                                                                                                                                                                                                                                                                                                                                                       63.0%;
78.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 ggtaagcagctgccattat 19
01-DEC-1997
                     ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26 GCTAACCAGCTGCTAATAT 8
                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 78.9 Matches 15; Conservative
                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS: ADDRESSEE: Rothwell,
                                                                                                                                                                                                                                                                                     ; MOLECULE TYPE: DNA
US-09-042-353-374
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-106-216-53/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: D
COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        ŏ
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E: Foley & Lardner
3000 K Street, N.W., Suite 500
                                                                                                           ZIP: 2007-5109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   61.0%;
82.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (202)672-5300
TELEFAX: (202)672-5399
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3 taagcagctgccattat 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 82.4
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (202)672-5399
TELEX: 904136
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5 TCATCAGGTGCCATTAT 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                Washington
    ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 14
US-08-472-802C-27
                                                                                               COUNTRY:
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APPLICANT: YARADA, Akiko
APPLICANT: KATO, Seishi
APPLICANT: KATO, Seishi
APPLICANT: KONDO, KIyosi
TITLE OF INVENTION: Gene Coding for Eicosapentaenoic Acid Synthesizing
TITLE OF INVENTION: Enzymes and Process for Production of Eiscosapentaenoic
TITLE OF INVENTION: Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
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                 Sequence 26, Application US/08482115B
Patent No. 5776679
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junli
APPLICANT: Feng, Junli
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Assays for the RNA Component of Human
TITLE OF INVENTION: Telomerase
NUMBER OF SEQUENCES: 40
CORRESPONDENCE: 40
CORRESPONDENCE: 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1; Length 25;
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                                                                                                                                                                                                                                                                                                                        CLILE: CCL.
STATE: CCL.
COMUTER: USA
ZIP: 94111-3834
COMPUTER READALE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1:0, Version #1:30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/482,115B
FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
FILING DATE: 07-JUL-1994
PRIOR APPLICATION DATE: 70-01L-1994
PRIOR APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
ATTORNEY/AGENT INFORMATION:
NAME: STORENIE 27-OCT-1994
ATTORNEY/AGENT INFORMATION:
NAME: STORENIE 27-OCT-1994
ATTORNEY/AGENT INFORMATION:
NAME: STORENIE 27-OCT-1994
TELECOMMUNICATION NUMBER: 32,944
REGISTRATION NUMBER: 32,944
REGISTRATION NUMBER: 32,544
REGISTRATION NUMBER: 32,544
RECEDENMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TRELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
"ENGTH: 25 base pairs
nucleic acide
nucleic acide
nucleic acide
nucleic acide
                                                                                                                                                                                                                                                                          ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 12.2;
Pred. No. 1e
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 61.0%;
Best Local Similarity 82.4%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 ggtaagcagctgccatt 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9 GGCCAGCAGCTGACATT 25
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CORRESPONDENCE ADDRESS:
US-08-482-115B-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-752-929-22
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0; , Gaps
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; Mismatches 3: ...
SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NAMBER: US/08/752,929 FILING DATE: 20-NOV-1996 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
LIP: 9411-3834
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/375,709
FILING DATE: 20-JAN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/178,251
FILING DATE: 14-MAY-1993
PRIOR APPLICATION NUMBER: US 08/178,251
FILING DATE: 15-MAY-1993
APPLICATION NUMBER: UP 4-147945
FILING DATE: 15-MAY-1992
ATTORNEY/AGBNT INFORMATION:
NAME: WEGNER, HAROLD C.
REGISTRATION NUMBER: 53466/150/AAOK
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 27, Application US/08472802C
Fatent No. 5958680
GENERAL INFORMATION:
APPLICANT: Villeponteau, Bryant
APPLICANT: Feng, Junii
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Mammalian Telomerase
NUMBER OF SEQUENCES: 44
CORRESPONDENCE ADDRESS:
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Gaps

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Score 12.2; DB 5; Length 41; Pred. No. 1.1e+03; 0; Mismatches 3; Indels
                                          REFERENCE DOCKET NUMBER: NEX40C/PCT TELECOMMUNICATION INFORMATION: TELEFONNE: (303) 793-3433
FILEFAX: 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Search completed: October 2, 2001, 16:03:49
Job time: 14593 sec
                   REGISTRATION NUMBER: 33,215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   61.0%;
82.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 gtaagcagctgccatta 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 61.0
Best Local Similarity 82.4
Matches 14; Conservative
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Pred. No. 1e+03;
0; Mismatches 3; Indels
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MEDIUM TYPE: Diskette, 3 1/2 diskette, 1.44 MB
COMPUTER: Diskette, 3 1/2 diskette, 1.44 MB
COPERATION GSYSTEM: MS-DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PSE CONTINUE PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/479,724
FILING DATE: 07-JUNE-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/472,256
FILING DATE: 07-JUNE-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/472,255
FILING DATE: 07-JUNE-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/477,829
FILING DATE: 07-JUNE-1995
PRIOR APPLICATION NUMBER: 08/477,829
FILING DATE: 07-JUNE-1995
ATTONNEY/AGENT INFORMATION:
AMAR: ABSENT APPLICATION NUMBER: 08/477,829
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Swanson & Bratschun, L.L.C.
STREET: 8400 E. Prentice Avenue, Suite 200
CITY: Englewood
STATE: Colorado
COUNTRY: USA
APPLICATION NUMBER: US/08/472,802C
ETLING DATE: 07-JUN-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/272,102
FILING DATE: 07-JUL-1994
FILING DATE: 07-JUL-1994
APPLICATION NUMBER: US 08/330,123
FILING DATE: 27-OCT-1994
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 15389-000820
TELEPHOME: (415) 576-0200
TELEPHOME: (415) 576-0300
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 61.0%;
Best Local Similarity 82.4%;
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 25 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; MOLECULE TYPE:
US-08-472-802C-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 80111
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116, Appl 19, Appl 19, Appl 4166, Ap 4167, Ap 4168, Ap 4170, Ap 4171, Ap 4172, Ap 4173, Ap 4174, Ap 4175, Ap 4176, Ap

Run on:

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Sequence 3994, Application US/09198452A

GENERAL INFORMATION:
APPLICANT: Griffals, R.
TITLE OF INVERTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragme
TITLE OF INVERTION: thereof and uses thereof, in particular for the diagnosis, pr
TITLE OF INVERTION: and treatment of infection
FILE REFERENCE: 9710-003-999
CURRENT APPLICATION NUMBER: US/09/198,452A
CURRENT FILING DATE: 1998-11-24
NUMBER OF SEQ ID NOS: 6849
SEQ ID NO 3994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      POLICES, Greg
VPENTION: PURIFICATION OF HIGHER ORDER TRANSCRIPTION
COMPLEXES FROM TRANSGENIC
NON-HUMAN ANIMALS
                                                        Sequence Seq
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ADDRESSEE: HELLER, EHRMAN, WHITE & MCAULIFFE
STREET: 1666 K Street, N.W., Suite 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 12.4; DB 6;
Pred. No. 1.1e+03;
); Mismatches 1;
US-09-866-108-2548
US-09-866-108-2549
US-09-866-108-2551
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US-09-866-108-2552
US-09-883-375-16
US-09-7330-5598-91
US-09-866-108-4166
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GENERAL INFORMATION:
APPLICANT: Kirschbaum, Bernd
Berglund, Erick
Meisterernst, Michael
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             82.7%;
92.9%;
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STATE: D.C.
COUNTRY: USA
ZIP: 20006
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  US-09-198-452A-3994
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Best Local Simi
Matches 13;
  LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1, Appli
Sequence 272, App
Sequence 272, App
Sequence 14, Appl
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Sequence 7, Appli
                                                                                                                                                                                   Search time 2173.58 Seconds (without alignments) 14.467 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Description
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1: /cgn2_6/ptodata/2/pna/PCT_NEW_COMB.seq:*

2: /cgn2_6/ptodata/2/pna/USO6_NEW_COMB.seq:*

3: /cgn2_6/ptodata/2/pna/USO8_NEW_COMB.seq:*

4: /cgn2_6/ptodata/2/pna/USO8_NEW_COMB.seq:*

5: /cgn2_6/ptodata/2/pna/USO9_NEW_COMB.seq:*

6: /cgn2_6/ptodata/2/pna/USO9_NEW_COMB.seq:*

7: /cgn2_6/ptodata/2/pna/USO9_NEW_COMB.seq:*

8: /cgn2_6/ptodata/2/pna/USO9_NEW_COMB.seq2:*
                            GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-198-452A-3994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1895189 seqs, 1048201267 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SUMMARIES
                                                                                                                                                                                   2001, 16:55:07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Maximum Match 100%
Listing first 45 summaries
                                                                                                                                   nucleic search, using sw model
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Minimum DB seq length: 0 Maximum DB seq length: 50
                                                                                                                                                                                         October
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                                                                                                                                   OM nucleic
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Database

Result No.

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Gaps

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TITLE OF INVENTION: Streptococcus pneumoniae Antigens and Vaccines NUMBER OF SEQUENCES: 452
CORRESPONDENCE ADDRESS:
ADDRESSEE: Human Genome Sciences, Inc.
STREET: 9410 Key West Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 272, Application US/09765272

GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: Streptococcus pneumoniae Antigens and Vaccines
NUMBER OF SEQUENCES: 452
CORRESPONDENCE ADDRESS:
ADDRESSE: Human Genome Sciences, Inc.
STREET: 9410 Key West Avenue
CITY: Rockville
STATE: Maryland
COUMTY: USA
ZIP: 20850
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                      STATE: MONTALLE
STATE: MASTALLE
COUNTRY: USA
ZIP: 20850
COMPUTER: READBLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage
COMPUTER: PH Vectra 486/33
OPERATING SYSTEM: MADOS Version 6.2
SOFTWARE: ASCII Text
CURRENT APPLICATION DATE:
RELING DATE: 30-Oct-1997
CLASSIFICATION NUMBER: US/09/536,784
FILING DATE: 30-Oct-1997
CLASSIFICATION NUMBER: 08/961,083
FILING DATE: OCT-30-1997
ATTORNEYAGENT INFORMATION:
NAME: Michelle S. Marks
REFERENCE DOCKET NUMBER: 41,971
FELECOMMUTICATION NUMBER: 41,971
FELECOMMUTICATION INFORMATION:
TELECOMMUTICATION INFORMATION:
TELECOMMUTICATION INFORMATION:
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COMPUTER: HP Vectra 486/33
COMPUTER: ASCII Text
CURRENT APPLICATION NUMBER: US/09/765,272
FILING DATE: 22-Jan-2001
CLASSIFICATION: <UNKNOWN>
PRIOR APPLICATION DATE:
APPLICATION NUMBER: 08/961,083
FILING DATE: <UNKNOWN>
FILING DATE: <UNKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 31;
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Pred. No. 4.6e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 272:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (301) 309-8504
TELEFAX: (301) 309-8512
INFORMATION FOR SEQ ID NO: 272:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS
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92.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 76.0
Best Local Similarity 92.3
Matches 12; Conservative
                                                                                                                          CITY: Rockville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3 gcttatcttcagt 15
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US-09-765-272-272
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GENERAL INFORMATION:
APPLICANT: Leiden, Jeffrey M.
APPLICANT: Leiden, Jeffrey M.
TITLE OF INVENTION: SYSTEM THAT REGULATES PROTEIN TRANSLATION
TITLE OF INVENTION: SYSTEM THAT REGULATES PROTEIN TRANSLATION
FILE REFERENCE: HUI-042
CURRENT APPLICATION NUMBER: US/09/815,980
CURRENT FILING DATE: 2001-03-23
PRIOR PHILNG DATE: 2000-03-23
NUMBER_OF SEQ ID NOS: 8
SOFTWARE: FastSEQ for Windows Version 4.0
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Pred. No. 2.6e+03;
); Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              78.7%; Score 11.8; DB 7; Length 21; 86.7%; Pred. No. 2.5e+03; ive 0; Mismatches 2; Indels
                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                       CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/849,243
FILING DATE: US/09/849,243
FILING DATE: US/09/849,243
FILING DATE: US/09/849,243
FILING DATE: US/08/849,243
FILING DATE: US/08/849,243
FILING DATE: US/08/849,243
FILING TAION NUMBER: 33,683
FEFENCE/DOCKET NUMBER: 38005-0148
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)912-2000
TELEPHONE: (202)912-2000
TELEPHONE: (202)912-2000
TELEPHONE: (202)912-2000
TELEPHONE: (202)912-2000
TELEPHONE: (202)912-2000
TELEPHONE: (202)912-2020
INFORMATION FOR SEQ. ID NO: 7:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LOCATION: 1..21
; SEQUENCE DESCRIPTION: SEQ ID NO: 7:
US-09-849-243-7
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GENERAL INFORMATION:
APPLICANT: Choi et. al.
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 21 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
MOLECULE TYPE: CDNA
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Best Local Similarity 86.7
Matches 13; Conservative
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LOCATION: 1..2
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; ORGANISM: Homo sapiens
US-09-815-980-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity
Matches 13; Conserv
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US-09-536-784-272
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US-09-815-980-1/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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Gaps

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                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Havukkala, Ilkka
TITLE OF INVENTION: Polynucleotide, Materials Incorporating
TITLE OF INVENTION: Them, and Methods for Using Them
FILE REFERENCE: 11000.1049AU
CURRENT APPLICATION NUMBER: US/09/724,750
CURRENT FILING DATE: 2000-11.28
PRIOR APPLICATION NUMBER: 60/171,431
PRIOR FILING DATE: 1999-12-21
NUMBER OF SEQ ID NOS: 25199
SOFTWARE: FASTSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Polynucleotide, Materials Incorporating TITLE OF INVENTION: Them, and Methods for Using Them FILE REFERENCE: 11000.104940 GCURENT APPLICATION NUMBER: US/09/724,750 CURRENT FILING DATE: 2000-11-28 PRIOR PPLICATION NUMBER: 06/171,431 PRIOR FILING DATE: 1999-12-21 NUMBER OF SEQ ID NOS: 25199 SOFFWARE: PastSEQ for Windows Version 4.0 SEQ ID NO 24046 LENGTH: 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  73.3%; Score 11; DB 7; Length 38; 100.0%; Pred. No. 8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 39;
                                                                                                                        Length 24;
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Pred. No. 8.1e+03;
0; Mismatches 0;
                                                                                                                        Score 11; DB 7; I
Pred. No. 7.4e+03;
                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                   MOLECULE TYPE: DNA
SEQUENCE DESCRIPTION: SEQ ID NO: 14:
US-09-717-438-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; Sequence 24046, Application US/09724750; GENERAL INFORMATION:
; APPLICANT: Havukkala, Ilkka
                                                                                                                                                                                                                                                                                                                                                     ; Sequence 14025, Application US/09724750; GENERAL INFORMATION:
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100.0%; Pre
0; }
                                                                                                                        73.3%; 8
                                                                                                                    Query Match 73.3
Best Local Similarity 100.
Matches 11; Conservative
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Best Local Similarity 100.
Matches 11; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
CORGANISM: Pinus radiata
US-09-724-750-14025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Pinus radiata
US-09-724-750-24046
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Best Local Similarity
Matches 11; Conserva
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27 ggcttatcttc 37
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 14025
LENGTH: 38
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Stemmer, Willem P.C.
Crameri, Andreas M.
TITLE OF INVENTION: Methods for Generating Polynucleotides
Having Desired Characteristics by Iterative Selection and
Recombination
                                                                                                                                                                                                                                                                                                                                                                                            0
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0
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ZIP: 94111-3834

COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30

CURRENT APPLICATION NUMBER: US/09/717,438
FILING DATE: 18-Nov-2000
CLASSIFCATION: <unimode classification = cunknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                            1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 70 CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
TOWNSENCE TOWNSENCE AND CLP
STREET: TWO Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Liebeschuetz, Joe
REGISTRATION NUMBER: 37,505
REFERENCE/DOCKET NUMBER: 018097-014612US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 08/621,859
FILING DATE: CURMOWN>
FILING DATE: CURMOWN>
FILING DATE: 30-607-1995
APPLICATION NUMBER: WO PCT/US95/02126
FILING DATE: 17-FEB-1995
APPLICATION NUMBER: US 08/198,431
FILING DATE: 17-FEB-1994
                                                                                                                                                                                                                                                                                                                                                 Score 11.4; DB 6;
Pred. No. 4.6e+03;
0; Mismatches 1;
                 NAME: Brookes, A. Anders
REGISTRATION NUMBER: 36,373
REFERENCE/DOCKET NUMBER: PB340P2
RELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8504
TELEFAX: (301) 309-8512
INFORMATION FOR SEQ ID NO: 272:
SEQUENCE CHARACTERISTICS:
LENGTH: 31 base pairs
                                                                                                                                                                                                                                              TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 272:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
INFORMATION FOR SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 14, Application US/09717438 GENERAL INFORMATION:
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ATTORNEY/AGENT INFORMATION
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LENGTH: 24 base pairs
                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                               76.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: California
                                                                                                                                                                                                                                                                                                                                                 Query Match 76.0
Best Local Similarity 92.3
Matches 12; Conservative
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US-09-717-438-14
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Sequence 1708, Application US/09818875

Sequence 1708, Application US/09818875

Sequence 1708, Application US/09818875

Sequence 1708, Application US/09818875

APPLICANT: Kmiec, Eric B.

APPLICANT: Gamper, Howard B.

APPLICANT: Rice, Michael C.

TITLE OF INVENTION: Stranded Oligonucleotides

FILE REFERENCE: Napro-4

CURRENT APPLICATION NUMBER: US/09/818,875

CURRENT APPLICATION NUMBER: US 60/192,176

PRIOR APPLICATION NUMBER: US 60/192,176

PRIOR APPLICATION NUMBER: US 60/192,179

PRIOR FILING DATE: 2000-03-27

PRIOR APPLICATION NUMBER: US 60/208,538

PRIOR FILING DATE: 2000-06-01

PRIOR FILING DATE: 2000-06-01

PRIOR FILING DATE: 2000-10-30

SOFTWARE FILING DATE: 2000-10-30

NUMBER OF SEQ ID NOS: 4385

SOFTWARE FILING DATE: 2000-10-30

SOFTWARE FILING DATE: 2000-10-30

SOFTWARE FILING DATE: 2000-10-30

LENGTH: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 10.8; DB 7;
Pred. No. 9.2e+03;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          72.0%; Score 10.8; DB 7;
85.7%; Pred. No. 9.2e+03;
ative 0; Mismatches 2;
CURRENT APPLICATION NUMBER: US/09/818,875
CURRENT FILING DATE: 2001-03-27
PRIOR APPLICATION NUMBER: US 60/192,176
PRIOR FILING DATE: 2000-03-27
PRIOR FILING DATE: 2000-03-27
PRIOR FILING DATE: 2000-03-27
PRIOR FILING DATE: 2000-06-01
PRIOR FILING DATE: 2000-06-01
PRIOR FILING DATE: 2000-06-01
PRIOR FILING DATE: 2000-10-30
PRIOR FILING DATE: 2000-10-30
SOFTWARE: FILING DATE: 2000-10-30
SOFTWARE: FILING DATE: 2000-10-30
NUMBER OF SEQ ID NOS: 4385
SOFTWARE: Friedman macro Napro4
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85.7%;
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APPLICANT: WADHWA, RENU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 72.0
Best Local Similarity 85.7
Matches 12; Conservative
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Best Local Similarity 85.7
Matches 12; Conservative
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                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
CRGANISM: Homo sapiens
US-09-818-875-1707
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US-09-818-875-1708
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                                                                             Sequence 13, Application US/09717438
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Stemmer, Willem P.C.
Crameri, Andreas M.
TITLE OF INVENTION: Methods for Generating Polynucleotides
Having Desired Characteristics by Iterative Selection and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 10
US-09-818-975-1707
US-09-818-975-1707
Sequence 1707, Application US/09818875
Sequence 1707, Application US/09818875
SEGUERAL INPORMATION:
APPLICANT: Kalec, Howard B.
APPLICANT: Rice, Michael C.
TITLE OF INVENTION: Targeted Chromosomal Genomic Alterations with Modified Single; TITLE OF INVENTION: Stranded Oligonucleotides
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                                                                                                                                                                                                                                                                           CORRESPONDER SEQUENCES:

ADDRESSE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco CITY: San Francisco COUNTRY: USA
CITY: San Francisco CONTRY: USA
LIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/09/717,438
FILING DATE: 18-Nov-2000
CLASSIFICATION NUMBER: US/08/517,874
FILING DATE: CURNOWN>
PRIOR APPLICATION NUMBER: US 08/537,874
FILING DATE: 30-OCT-1995
APPLICATION NUMBER: US 08/537,874
FILING DATE: 17-FEB-1995
APPLICATION NUMBER: US 08/537,874
FILING DATE: 17-FEB-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               73.3%; Score 11; DB 7; Length 41; 100.0%; Pred. No. 8.1e+03; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Liebeschuetz, Joe
REGISTRATION NUMBER: 37,505
REFERENCE/DOCKET NUMBER: 018097-014612US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; TOPOLOGY: 11near; MOLECULE TYPE: DNA SEQUENCE DESCRIPTION: SEQ ID NO: 13: US-09-717-438-13
                                                                                                                                                                                                                                        Recombination
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0200
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
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STRANDEDNESS: single
                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 100.
Matches 11; Conservative
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OTHER INFORMATION: nucleic acid probe
NAME/KEY: misc_feature
LOCATION: (1)..(1)
OTHER INFORMATION: (1)..(2)
COTHER INFORMATION: (1)..(23)
OTHER INFORMATION: (1)..(23)
OTHER INFORMATION: probe
NAME/KEY: misc_feature
LOCATION: (2)..(23)
OTHER INFORMATION: 3.0-(7.coumariny1).glycerol between position 22 and 23
US-09-379-888B-8
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APPLICANT: Albagli, David
APPLICANT: Albagli, David
APPLICANT: Wood, Michael L.
APPLICANT: Wood, Atta, Reuel B.
APPLICANT: Cheng, Peter C.
TITLE OF INVENTION: Assays Using Crosslinkable Immobilized Nucleic Acids
FILE REPERENCE: 14251-0006 crp
CURRENT APPLICATION NUMBER: US/09/379,888B
CURRENT FILING DATE: 1999-08-23
NUMBER OF SEQ ID NOS: 21
SOFTWARE: Patentin version 3.0
SQTUARE: Patentin version 3.0
                                                                                                                                                                                               Gaps
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                                                                                                                                                    Length 21;
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Pred. No. 9.6e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                             Indels
                                                                                                                                                  Score 10.8; DB 7;
Pred. No. 9.5e+03;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Search completed: October 2, 2001, 16:55:07 Job time: 17670 sec
                                                                 ; OTHER INFORMATION: Synthetic Sequence US-09-920-313-140
                                                                                                                                                                                                                                                                                                                                                                           Sequence 8, Application US/09379888B GENERAL INFORMATION:
                           ORGANISM: Artificial Sequence
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85.7%;
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85.7%;
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Best Local Similarity 85.7
Matches 12; Conservative
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Best Local Similarity 85.7
Matches 12; Conservative
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GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragments
TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, preve
TITLE OF INVENTION: and treatment of infection
FILE REPERBNCE: 9710-003-999
CURRENT FILING DATE: 1998-11-24
NUMBER OF SEQ ID NOS: 6849
SEQ ID NO 1913
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APPLICANT: KAUL, SUNIL C.
APPLICANT: MITSUI, YOUJI
TITLE OF INVENTION: TESTIS-SPECIFIC DIFFERENTIATION-REGULATORY FACTOR
TITLE REFERENCE: 084335/0127
CURRENT APPLICATION NUMBER: US/09/743,237
CURRENT FILING DATE: 2001-06-04
PRIOR FILING DATE: 1999-07-16
NUMBER OF SEQ ID NOS: 24
SOFTWARE: PATENTIN VET. 2.1
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; OTHER INFORMATION: Description of Artificial Sequence: Primer
US-09-743-237-16
                                                                                                                                                                                                                                                                                                                                                                                              Length 20;
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GENERAL INFORMATION:
APPLICANT: Bratzler, Robert L.
APPLICANT: Petersen, Deanna M.
TITLE OF INVENTION: Nucleic Acids for the Prevention and
TITLE OF INVENTION: Treatment of Gastric Ulcers
FILE REFERENCE: C1037/7019 (HCL/MAT)
CURRENT APPLICATION NUMBER: US/09/920,313
CURRENT FILING DATE: 2001-08-01
PRIOR FILING DATE: 2001-08-08
NUMBER OF SEQ ID NOS: 148
SOFTWARE: FastSEQ for Windows Version 3.0
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Pred. No. 9.4e+03;
0; Mismatches 2;
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Pred. No. 9.4e+03;
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ORGANISM: Chlamydia pneumoniae
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85.7%;
                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Artificial Sequence
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19 GGCTTGTCTTGAGT 6
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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                                                                                                                           APPLICANT: IS15 Pharmaceuticals, Inc.
APPLICANT: Monia, Brett P.
APPLICANT: Gardé, William A.
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0476
CURRENT APPLICATION NUMBER: PCT/US00/18999
CURRENT APPLICATION NUMBER: PCT/US00/18999
FILE REPERIORE: 1999-08-19
FILE APPLICATION NUMBER: 09/377,310
PRIOR FILING DATE: 1999-08-19
NUMBER OF EXQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 33
LENGHH: 15
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APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT FILING NUMBER: US/09/377,310A
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 33
LENGTH: 15
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100.0%; Pred. No. 3.2e+02;
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ALIGNMENTS
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                                                                                            Sequence 33, Application PC/TUS0018999 GENERAL INFORMATION:
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US-09-757-100B-33
; Sequence 33, Application US/09757100B
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Sequence 33, Application US/09377310A
; GENERAL INFORMATION:
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Best Local Similarity 100.
Matches 15; Conservative
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Best Local Similarity
Matches 15; Conserva
                                                                            PCT-US00-18999-33
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APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Garade, William A.
TAPLICANT: Geral Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REPERENCE: ISPH-0476
CURRENT APPLICATION NUMBER: PCT/US00/18999
CURRENT APPLICATION NUMBER: PCT/US00/18999
FRIOR APPLICATION NUMBER: 09/377,310
PRIOR APPLICATION NUMBER: 09/377,310
PRIOR FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 13
LENGTH: 20
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           APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Expression
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0533
CURRENT APPLICATION NUMBER: 02/09/757,100B
CURRENT APPLICATION NUMBER: 09/377,310
PRIOR APPLICATION NUMBER: PCT/US00/18999
PRIOR FILING DATE: 1999-08-19
PRIOR FILING DATE: 2000-07-13
NUMBER OF SEQ ID NOS: 44
SEQ ID NOS: 44
LENGTH: 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 100.0%; Score 15; DB 29; Best Local Similarity 100.0%; Pred. No. 3.2e+02; Matches 15; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 15; DB 1; 100.0%; Pred. No. 3.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: antisense sequence US-09-757-1008-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: antisense sequence PCT-USO0-18999-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 4
CT-1050-18999-13
Sequence 13, Application PC/TUS0018999
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 15; Conservative
APPLICANT: Monia, Brett P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Hasegawa, Osamu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 tggcttatcttcagt 15
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Query Match
Best Local Similarity
Matches 14; Conserv
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Best Local Similarity
Matches 15; Conserv
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US-60-233-166-354400
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GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Gaarde, William A.

TITLE OF INVENTION: Expression

FILE REFERENCE: ISPH-0389

CURRENT APPLICATION UNBER: US/09/377,310A

CURRENT FILING DATE: 1999-08-19

NUMBER OF SEQ ID NOS: 43

SOFTWARE: Patentin Ver. 2.0

LENGTH: 20
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GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
APPLICANT: Generate, William A.
APPLICANT: Nero, Pamela S.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0533
CURRENT FILING NUMBER: US/09/757,100B
PRIOR APPLICATION NUMBER: 09/377,310
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                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: Description of Artificial Sequence: US-09-050-392-15
                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 15; DB 14; 100.0%; Pred. No. 3.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
100.0%; Score 15; DB 17;
Best Local Similarity 100.0%; Pred. No. 3.4e+02;
Matches 15; Conservative 0; Mismatches 0;
                               APPLICANT: Indra, Yuri
TILE OF INVENTION: Cellulose Synthase Gene
FILE REFERENCE: 2011-0017
CURRENT APPLICATION NUMBER: US/09/050,392A
CURRENT FILING DATE: 1998-03-30
EARLIER FILING DATE: 1997-04-01
NUMBER OF SEQ ID NOS: 24
SOFTWARE: Patentin Ver: 2.0
SEQ ID NO 15
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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APPLICANT: Aotsuka, Satoshi
APPLICANT: Hayashi, Takahisa
                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 tggcttatcttcagt 15
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APPLICANT: Clumakov, Ilya
APPLICANT: Chumakov, Ilya
APPLICANT: Chumakov, Ilya
ATITLE OF INVENTION: Biallelic markers for use in constructing a high density...
FILE REPERENCE: GENSET.020cp1
CURRENT APPLICATION NUMBER: US/09/422,978
CURRENT APPLICATION NUMBER: US 09/298,850
EARLIER FILING DATE: 1999-04-21
EARLIER FILING DATE: 1999-04-21
EARLIER FILING DATE: 1998-11-23
EARLIER FILING DATE: 1998-11-23
EARLIER FILING DATE: 1998-04-21
NUMBER OF SEQ ID NOS: 11796
SEQ ID NO 3990
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                                                                                                                                                                                                                                                                          Length 20;
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APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of the Rat
TITLE OF INVENTION: Genome
                                                                                                                                                                                                                                                                                                                 Indels
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Pred. No. 2.8e+03;
0; Mismatches 1;
                                                                                                                                                                                                                                                                        100.0%; Score 15; DB 29; 100.0%; Pred. No. 3.4e+02;
                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: 3112
CURRENT APPLICATION NUMBER: US/60/233,166
CURRENT FILING DATE: 2000-10-24
NUMBER OF SEQ ID NOS: 420907
SOFTWARE: FastSEQ for Windows Version 4.0
LENGTH: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank U65486
PRIOR APPLICATION NUMBER: PCT/US00/18999
PRIOR FILING DATE: 2000-07-13
NUMBER OF SEQ ID NOS: 44
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 13
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 354400, Application US/60233166 GENERAL INFORMATION:
                                                                                                                                                                                   ; OTHER INFORMATION: antisense sequence US-09-757-100B-13
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                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence
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93.3%;
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                                                                                                                                                                                                                                                                                                                 15; Conservative
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APPLICANT: Huang, Shi
TITLE OF INVENTION: Retinoblastoma Protein - Interacting
TITLE OF INVENTION: Zinc Finger Proteins
NUMBER OF SEQUENCES: 106
CORRESPONDENCE ADDRESS:
  Retinoblastoma Protein - Interacting
                                                                                                                                                                                                                                                                                       SOFTWARE PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/516,859
FILING DATE: 18 A.0G-1995
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/POCKET UNBER: P-LA 1776
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
TELEFEX: (619) 535-9049
: INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
                                                                                     STREET: 4370 La Jolla Village Drive, Suite 700 CIT: San Diego STATE: California COUNTR: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/528,706
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Campbell & Flores LLP
STREET: 4370 La Jolla Village Drive, Suite 700
CITY: San Diego
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              82.7%; Score 12.4; DB 9; 92.9%; Pred. No. 1.1e+04;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No. 1.16; Mismatches
                      Zinc Finger Proteins
106
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/516,859
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/292,683
                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-528-706-11/c; Sequence 11, Application US/09528706; GENERAL INFORMATION:
                                                                                   Campbell and Flores
                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27 base pairs
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                                         NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 tggcttatcttcag 14
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TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
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Best Local Similarity
Matches 13; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE
                                                                                 ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        92122
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 12
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                                                         FEATURE:
NAME/KEY: primer_bind
LOCATION: 1.18
COTHER INFORMATION: upstream amplification primer 99-12619 for SEQ 56, US-09-422-978-3990
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                                                                                                                                                                                                          Length 18;
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                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Huang, Shi
TITLE OF INVENTION: Nucleic Acid Molecules Encoding RIZ
TITLE OF INVENTION: Protein
NUMBER OF SEQUENCES: 90
CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell and Flores
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION STSTEM: CLOSS/MS-LOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/292,683
FILING DATE: 18-Aug-1994
CLASSIFICATION: 514
ATORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REGISTRATION NUMBER: P-LJ 1004
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION OF $155-9001
TELEFAK: (619) 535-9001
TELEFAK: (619) 535-8949
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Campbell and Flores
STREET: 4370 La Jolla Village Drive, Suite 700
CITY: San Diego
STATE: California
                                                                                                                                                                                                        82.7%; Score 12.4; DB 18; 92.9%; Pred. No. 1e+04; ive 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
ZIP: 92122
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 11, Application US/08292683 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 11, Application US/08516859
GENERAL INFORMATION:
APPLICANT: Huang, Shi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         82.7%;
92.9%;
                                                                                                                                                                                                      Query Match 82.7
Best Local Similarity 92.9
Matches 13; Conservative
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Best Local Similarity 92.9
Matches 13; Conservative
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                                                                                                                                                                                                                                                                                            2 ggcttatcttcagt 15
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                   TYPE: DNA
ORGANISM: Homo Sapiens
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                                                                                                                                                                                                                                                                                                                    18 GCCTTATTTCAGT
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US-08-292-683-11/c
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LENGTH: 18
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Gaps

Length 27;

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APPLICANT: Cohen, Daniel
APPLICANT: Cohen, Marta
APPLICANT: Blumenfeld, Marta
APPLICANT: Chumakov, 11ya
APPLICANT: Abderrahim, Hadi
APPLICANT: Dufaure-Gare, Isabelle
TITLE OF INVENTION: BIALLELIC MARKER MAPS FOR USE IN CONSTRUCTING A HIGH DENSITY.
FILER REFERENCE: 84.US1.PRO
CURRENT APPLICATION NUMBER: US/60/216,745
CURRENT FILING DATE: 2000-06-30
NUMBER OF SEQ ID NOS: 13665
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; NAME/KEY: primer_bind
; LOCATION: 1..18
; OTHER INFORMATION: upstream amplification primer 99-46755 for SEQ 3132,
US-60-216-745-7663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Tang, Jianming
APPLICANT: Tang, Jianming
APPLICANT: Raslow, Richard A.
TITLE OF INVENTION: Methods of Surveying For CC (Beta) Chemokine
TITLE OF INVENTION: Receptor Variants and Their Association With HIV-1
TITLE OF INVENTION: Transmission and/or Disease Progression
TITLE OF INVENTION: Transmission and/or Disease Progression
CURRENT APPLICATION NUMBER: US/09/638,509C
CURRENT APPLICATION NUMBER: 60/148,530
PRIOR PELING DATE: 1999-08-11
PRIOR PELING DATE: 1999-08-12
NUMBER OF SEQ ID NOS: 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: CCR5-PM6G, primer used for typing major CTER INFORMATION: polymorphism in CCR2b, CCR5 and the CCR5 downstream OCTER INFORMATION: promoter region US-09-638-509C-18
                                                                                                          Gaps
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                                                                   Length 27;
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                                                                                                          Indels
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                                                               Score 12.4; DB 22
Pred. No. 1.1e+04;
); Mismatches 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 12;
Pred. No.
                                                                                                                                                                                                                                                                                          Sequence 7663, Application US/60216745; GENERAL INFORMATION:
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28.09-618/C
; Sequence 18, Application US/09638509C
; GENERAL INFORMATION:
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100.0%; Pre
0;
                                                               82.7%;
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                                                            Query Match 82.7
Best Local Similarity 92.9
Matches 13; Conservative
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                                                                                                                                                                     27 TGGCTTTTCTTCAG 14
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ORGANISM: Homo Sapiens
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Best Local Similarity
Matches 12; Conserva
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                                                                                                                                               1 tggcttatcttcag
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SEQ ID NO 7663
  US-09-586-472-11
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LENGTH: 19
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COMPUTER READABLE FORM:

MEDIUW TYPE: Floppy disk

COMPUTER: IBM PC Compatible

COMPUTER: IBM PC COMPATIBLE

COMPUTER: DEAD FOLDS/MS-DOS

SOFTWARE: PATENTIN RE-BASE #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/586,472

FILING DATE: 01-Jun-2000

CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 09/528,706

FILING DATE: 17-MAR-1995

APPLICATION NUMBER: US 08/316,859

FILING DATE: 06-MAR-1995

APPLICATION NUMBER: US 08/39,411

FILING DATE: 18-AUG-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 11, Application US/09586472
GENERAL INFORMATION:
APPLICANT: Huang, Shi
TITLE OF INVENTION: Retinoblastoma Protein - Interac
NUMBER OF SEQUENCES: 106
CORRESPONDENCE ADDRESS:
ADDRESSE: Campbell & Flores LLP
STREET: 4370 La Jolla Village Drive, Sulte 700
                                                                                                                                                                                                                                                                                                                                                         DB 19;
                                                                                                                                                                                                                                                                                                                                                                                                   1;
                                                                                                                                                                                                                                                                                                                                                                         1.1e+04;
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                                                                                                                                                                                                                                                                                                                                                         Score 12.4;
Pred. No. 1.1
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SEQUENCE DESCRIPTION: SEQ ID NO: 11:
                                                                                P-LJ 1776
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Campbell, Cathryn A. REGISTRATION NUMBER: 31,815
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (619) 535-9001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION
                                NAME: Cambell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFRENCE/DOCKET NUMBER: P-LJ
TELECOMMULCATION INFORMATION:
TELEPHONE: (619) 535-9001
TELEPHONE: (619) 535-8949
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 27 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (619) 535-8949 INFORMATION FOR SEQ ID NO: 11:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
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92.9%;
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FILING DATE: 18-AUG-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: San Diego
STATE: California
                   INFORMATION
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Best Local Similarity 92.9
Matches 13; Conservative
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-09-586-472-11/c
                                                                                                                                                                                                                                                                       ; TOPOLOGY:
US-09-528-706-11
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0; Gaps
Query Match

80.0%; Score 12; DB 24; Length 19;
Best Local Similarity 100.0%; Pred. No. 1.7e+04;
Matches 12; Conservative 0; Mismatches 0; Indels
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Search completed: October 2, 2001, 21:50:22 Job time: 24535 sec

1624, Ap 5614, Ap 42408, A 16237, A 30106, A

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Run

Sequence:

Searched:

Database

Result Š

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Sequence 122, Application US/09672217

GENERAL INFORMATION:
APPLICANT: FELL, JACK
APPLICANT: BELL, JACK
APPLICANT: BIAZ, MARA
APPLICANT: DIAZ, MARA
APPLICANT: DIAZ, MARA
APPLICANT: DIAZ, MARA
CURRENT MCABE, MEADE
FILE REPERENCE: 086222/0143
CURRENT APPLICATION NUMBER: US/09/672,217
CURRENT FILING DATE: 1909-09-29
PRIOR PILING DATE: 1909-09-29
NUMBER OF SEQ ID NOS: 301
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: Description of Artificial Sequence: Primer/Probe US-09-672-217-122
                              Sequence Seq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
Sequence
Sequence
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APPLICANT: FELL, JACK
APPLICANT: DIAZ, MARA
APPLICANT: DIAZ, MARA
APPLICANT: DIAZ, MARA
APPLICANT: DIAZ, MARA
TITLE OF INVENTION: METHOD OF IDENTIFYING PATHOGENIC CRYPTOCOCCI
FILE REFERENCE: 086222/0143
CURRENT PAPLICATION NUMBER: US/09/672,217
CURRENT FILING DATE: 2000-09-29
PRIOR PALING DATE: 1999-09-29
NUMBER OF SEQ ID NOS: 301
SOFTWARE: PALENTIN Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       82.7%; Score 12.4; DB 7; Length 33; 92.9%; Pred. No. 1.1e+03; ive 0; Mismatches 1; Indels
US-09-866-108-12612
US-09-866-108-12613
US-09-866-108-12614
US-09-866-108-12616
US-09-866-108-12616
US-09-866-108-12617
US-09-817-1624
US-09-817-819-5614
US-09-817-819-5614
US-09-811-819-5614
US-09-811-819-5614
US-09-811-819-511
US-09-811-918-51
US-09-841-353A-10
US-09-648-103B-51
US-09-648-103B-51
US-09-648-103B-51
US-09-648-103B-51
US-09-648-1332-10744
US-09-481-375-29
US-09-486-108-1321
US-09-866-108-1321
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ORGANISM: Artificial Sequence
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Best Local Similarity 92.9
Matches 13; Conservative
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   RESULT 2
US-09-672-217-223
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Sequence 294, App
Sequence 7712, Ap
Sequence 7714, Ap
Sequence 7714, Ap
Sequence 7715, Ap
Sequence 7715, Ap
Sequence 165, App
Sequence 165, App
Sequence 166, App
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 122, App
Sequence 223, App
Sequence 65, Appl
Sequence 1705, Ap
Sequence 10, Appl
Sequence 4, Appli
                                                                                                                                                                                                  Search time 2173.58 Seconds (without alignments) 14.467 Million cell updates/sec
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Sequence 1
Sequence 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /cgn2_6/ptodata/2/pna/PCT_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/US06_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/US08_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/US08_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/US09_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/US09_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/US09_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/US09_NEW_COMB.seq:*
/cgn2_6/ptodata/2/pna/US09_NEW_COMB.seq:*
                           GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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US-09-87-281-65
US-09-801-274-1705
US-09-101-274-1705
US-09-406-643-293
US-09-406-643-293
US-09-866-108-7712
US-09-866-108-7712
US-09-866-108-7714
US-09-866-108-7715
US-09-866-108-7715
US-09-866-108-7716
US-09-866-108-7716
US-09-866-108-7716
US-09-866-108-7716
US-09-927-796-168
US-09-927-796-168
US-09-927-796-168
US-09-927-796-168
US-09-966-108-12604
US-09-866-108-12604
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US-09-866-108-12605
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US-09-866-108-12609
US-09-866-108-12610
US-09-866-108-12611
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1895189 seqs, 1048201267 residues
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                                                                                                                                                                                                  October 2, 2001, 16:55:07
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Listing first 45 summaries
                                                                                                                                             - nucleic search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                            IDENTITY_NUC
Gapop 10.0 , Gapext 1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pending_Patents_NA_New:*
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15
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Match
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Score 11.4; DB 6; Pred. No. 4.3e+03; 0; Mismatches 1;

76.0%; 92.3%;

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; Sequence 10, Application US/09794384A; GENERAL INFORMATION: APPLICANT: Dellaporta, Stephen L.
                                                                                                      Best Local Similarity 92.3
Matches 12; Conservative
                                                                                                                                                                 3 cccaggtgagtct 15
||||| ||||||||||||
14 CCCAGCTGAGTCT 2
  ; ORGANISM: Homo sapiens
US-09-801-274-1705
                                                                                                                                                                                                                                                                           RESULT 5
US-09-794-384A-10
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LENGTH: 27
                                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        ## APPLICANT: B. I. du Pont de Nemours and Company
## APPLICANT: B. I. du Pont de Nemours and Company
## TITLE OF INVENTION: Nucleic Acid Sequences Encoding Isoflavone Synthase
## FILE REFERENCE: BB1339 PCT
## CURRENT APPLICATION NUMBER: US/09/857,581
## CURRENT FILING DATE: 2001-06-05
## PRIOR APPLICATION NUMBER: 60/117,769
## PRIOR PILING DATE: 1999-01-27
## PRIOR PILING DATE: 1999-01-27
## PRIOR PILING DATE: 1999-07-20
## PRIOR APPLICATION NUMBER: 60/144,783
## PRIOR PILING DATE: 1999-07-20
## PRIOR PILING DATE: 1999-07-20
## PRIOR PILING DATE: 1999-02-24
## NUMBER OF SEO ID NOS: 66
## SEO ID NO 65
## LENTH: 25
## LENTH: 25
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0
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                                     TYPE: DNA
ONCHINEM: Artificial Sequence
FEATURE:
COTHER INFORMATION: Description of Artificial Sequence: Primer/Probe
US-09-672-217-223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: Description of Artificial Sequence: PCR PRIMER US-09-857-581-65
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                                                                                                                                                                                            Length 33;
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APPLICANT: Cargill, Michele
APPLICANT: Lander, Eric S.
TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
FILE REFERENCE: 2825.2009-001
CURRENT APPLICATION NUMBER: US/09/801,274
CURRENT FILING DATE: 2001-03-07
PRIOR APPLICATION NUMBER: US 60/187,510
PRIOR PILING DATE: 2000-03-07
PRIOR PELING DATE: 2000-05-22
NUMBER OF SEQ ID NOS: 1802
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1705
TYPE: DNA
                                                                                                                                                                                                                                   1; Indels
                                                                                                                                                                                     Score 12.4; DB 7;
Pred. No. 1.1e+03;
0; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                                             RESULT 3
US-09-857-581-65/c
; Sequence 65, Application US/09857581
; GENERAL INFORMATION:
                                                                                                                                                                                   82.7%;
92.9%;
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Best Local Similarity 92.3%;
Matches 12; Conservative
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Best Local Similarity 92.9
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                2 acccaggtgagtct 15
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US-09-801-274-1705/c
SEQ ID NO 223
               LENGTH: 33
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APPLICANT: Moreover, and a particular a particul
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APPLICANT: Ludwig, Janos
APPLICANT: Ludwig, Janos
APPLICANT: Sproat, Brian
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: Compositions Having RNA Cleaving Activity
FILE REFERENCE: MBHB00-874-A (237/197)
CURRENT APPLICATION NUMBER: US/09/406,643
CURRENT FILING DATE: 1999-09-27
PRIOR FILING DATE: 1997-06-19
PRIOR PILING DATE: 1997-06-19
PRIOR FILING DATE: 1997-06-19
NUMBER OF SEQ ID NOS: 2606
SOFTWARE: PatentIn version 3.0
SEQ ID NO 4
LENGTH: 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 10.8; DB 7;
Pred. No. 9.7e+03;
); Mismatches 2;
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85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 72.0
Best Local Similarity 85.7
Matches 12; Conservative
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SEGNERAL INFORMATION:

APPLICANT: 01, Vonggang
APPLICANT: 02, Vonggang
APPLICANT: 03, Vonggang
APPLICANT: 04, VONGSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AROMICAS: US/09/866, 108
CURRENT FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-09-27
PRIOR PELICATION NUMBER: 05 60/236, 359
PRIOR FILING DATE: 2001-00-3
PRIOR PELICATION NUMBER: PCT/USO1/0066
PRIOR PELICATION NUMBER: PCT/USO1/0066
PRIOR APPLICATION NUMBER: PCT/USO1/0066
PRIOR PRIOR APPLICATION NUMBER: PCT/USO1/0066
PRIOR PLILNG DATE: 2001-01-30
PRIOR PLING DATE: 2001-01-30
PRIOR PLING DATE: 2001-01-30
PRIOR DA
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APPLICANT: GI, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
                                                                                                                                                  Sequence 7712, Application US/09866108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11; Conservative
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; ORGANISM: Homo sapiens
US-09-866-108-7712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3 cccaggtgagtc 14
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      13 CCAGGTGGGTCT
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                                                                                                                                                                                                                                                                                  Sequence 293, Application US/09406643
GENERAL INFORMATION:
APPLICANT: Ribozyue Pharmaceuticals, Inc.
APPLICANT: Sproat, Brian
APPLICANT: Beigelman, Leonid
TITLE OF INVENTION: Compositions Having RNA Cleaving Activity
FILE REFERENCE: MBHB00-874-A (237/197)
CURRENT FILING DATE: 1999-09-27
PRIOR APPLICATION NUMBER: US 08/879,078
PRIOR FILING DATE: 1997-06-19
NUMBER OF SEQ ID NOS: 2606
SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: LUdwig, Janos
APPLICANT: Brian
APPLICANT: Brian
APPLICANT: Brian
APPLICANT: Brian
TITLE OF INVENTION: Compositions Having RNA Cleaving Activity
FILE REFERENCE: MBHB00-874-A (237/197)
CURRENT PPLICATION NUMBER: US/09/406,643
CURRENT FILING DATE: 1999-09-27
PRIOR RILING DATE: 1997-06-19
PRIOR PLILING DATE: 1997-06-19
PRIOR FILING DATE: 1997-06-19
PRIOR FILING DATE: US 08/878,640
PRIOR FILING DATE: US 08/878,640
PRIOR FILING DATE: US 08/878,640
SPRIOR FILING DATE: US 08/878,640
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Pred. No. 1.6e+04;
0; Mismatches 1; Indels
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                              1; Indels
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Pred. No. 1.6e+04;
      Pred. No. 1.6e+04;
                              Mismatches
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GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
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Best Local Similarity 91.7%;
Matches 11; Conservative
Best Local Similarity 91.7 Matches 11; Conservative
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Best Local Similarity 91.7
Matches 11; Conservative
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ORGANISM: Homo sapiens
US-09-406-643-294
                                                                                  4 ccaggtgagtct 15
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Pred. No. 1.6e+04;
); Mismatches 1;
                  PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR PLING DATE: 2000-09-21
PRIOR PLING DATE: 2000-09-21
PRIOR PLING DATE: 2000-09-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Aeomica Sequence Listing Engine
SEQ ID NO 7714
LENGTH: 17
FILING DATE: 2000-10-04
APPLICATION NUMBER: US 60/236,359
FILING DATE: 2000-09-27
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Best Local Similarity 91.7%;
Matches 11; Conservative
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APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 15752
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; ORGANISM: Homo sapiens
US-09-866-108-7714
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                                        APPLICANT: GHEN, Wensheng
APPLICANT: GENANON, MAIK
ITILE REPERENCE: ADOMICA-7
CURRENT PETLING DATE: 2001-05-25
CURRENT PILLING DATE: 2001-05-25
PRIOR PILLING DATE: 2000-05-26
PRIOR PLILING DATE: 2000-05-27
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR PPLICATION NUMBER: DC 000-05-27
PRIOR APPLICATION NUMBER: PCT/USO1/00665
PRIOR PPLICATION NUMBER: PCT/USO1/00667
PRIOR PLILING DATE: 2001-01-30
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Pred. No. 1.6e+04;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Aeomica Sequence Listing Engine
SEQ ID NO 7713
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 91.7
Matches 11; Conservative
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; ORGANISM: Homo sapiens
US-09-866-108-7713
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5 cctaggtgagtc 16
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APPLICANT: RANKEL DATA R. APPLICANT: RANKEL DATA R. APPLICANT: CHEN, Wensheng
APPLICANT: CHEN, Wensheng
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, MAY.
TITLE OF INVENTION: WYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
FULCE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
PRIOR PRILING DATE: 2000-05-26
PRIOR PRILING DATE: 2000-05-26
PRIOR PRILING DATE: 2000-10-04
PRIOR PLILING DATE: 2000-10-04
                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/USO1/00666
PRIOR FILING DATE: 2001-01-30
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APPLICANT: GV. YIZHONG
APPLICANT: GV. YIZHONG
APPLICANT: JI, YONGGANG
APPLICANT: PRNK, Sharron G.
APPLICANT: FANK, David K.
APPLICANT: RANK, David K.
APPLICANT: RANK, David K.
APPLICANT: RANK, David R.
APPLICANT: SHANNON, MANAL LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION, NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR FILING DATE: 2000-05-26

Application US/09866108

; Sequence 7714,

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APPLICANT: HANKEL, DAVIG R.
APPLICANT: HANKEL, DAVIG R.
APPLICANT: GIEN, Wensheng
APPLICANT: GIEN, Wensheng
APPLICANT: GIEN, Wensheng
APPLICANT: SHANNON, MAT.
TITLE OF INVENTION: WOOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
ITTLE REFERENCE: AECMICA-7
CURRENT APPLICATION NUMBER: US/02/9/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/230,456
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR APPLICATION NUMBER: DC 60/236,359
PRIOR APPLICATION NUMBER: PCT/USO1/00666
PRIOR APPLICATION NUMBER: PCT/USO1/00666
PRIOR PLING DATE: 2001-01-30
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Pred. No. 1.6e+04;
); Mismatches 1
**RIOK FILING DATE: 2001-01-30

**PRIOR APPLICATION NUMBER: PCT/USO1/00670

**PRIOR FILING DATE: 2001-01-30

**PRIOR PRILING DATE: 2000-09-21

**PRIOR APPLICATION NUMBER: US 60/234,687

**PRIOR APPLICATION NUMBER: US 60/266,860

**PRIOR APPLICATION NUMBER: US 60/266,860

**PRIOR FILING DATE: 2001-02-05

**NUMBER: ABOMICA ABOMICA SEQUENCE LISTING ENGINE

**SEQ ID NO 7716

**LENGTH: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 2000-09-21
APPLICATION NUMBER: US 60/266,860
FILING DATE: 2001-02-05
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; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 69.3
Best Local Similarity 91.7
Matches 11; Conservative
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; ORGANISM: Homo sapiens
US-09-866-108-7716
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APPLICANT: J1, Yonggang
APPLICANT: J1, Yonggang
APPLICANT: J1, Yonggang
APPLICANT: J1, Yonggang
APPLICANT: BNN, Sharron G.
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: RANK, David R.
APPLICANT: SHANNON, Mark
TILLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REPREBACE: SOOD-05-26
PRIOR APPLICATION NUMBER: US 60/207, 456
PRIOR PLILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-06-46
PRIOR PRIOR APPLICATION NUMBER: US 60/236, 359
PRIOR PLICATION NUMBER: US 60/236, 359
PRIOR PLICATION NUMBER: PCT/US01/0066
PRIOR PLICATION NUMBER: PCT/US01/0066
PRIOR PLILING DATE: 2001-01-30
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Pred. No. 1.6e+04;
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              SOFTWARE: Acomica Sequence Listing Engine SEQ ID NO 7715
APPLICATION NUMBER: PCT/US01/00669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: PCT/US01/00663 PRIOR FILING DATE: 2001-01-30 PRIOR APPLICATION NUMBER: PCT/US01/00662 PRIOR FILING DATE: 2001-01-30
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Sequence 7716, Application US/09866108
GENERAL INFORMATION:
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91.7%;
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; ORGANISM: Homo sapiens
US-09-866-108-7715
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Best Local Similarity
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R APPLICATION NUMBER: 60/133459

R FILING DATE: 1999-05-11

R APPLICATION NUMBER: 60/140650

R APPLICATION NUMBER: 60/140653

R APPLICATION NUMBER: 60/14053

R APPLICATION NUMBER: 60/144758

R FILING DATE: 1999-07-20

R FILING DATE: 1999-07-26

R APPLICATION NUMBER: 60/14629

R FILING DATE: 1999-07-26

R APPLICATION NUMBER: 60/14622

R APPLICATION NUMBER: 60/146395

R FILING DATE: 1999-07-28

R APPLICATION NUMBER: 60/146395

R FILING DATE: 1999-08-17
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RAPLICATION NUMBER: 09/143068

RETLING DATE: 1998-08-28

RETLING DATE: 1998-08-28

RETLING DATE: 1998-08-28

RETLING DATE: 1998-09-18

RETLING DATE: 1998-09-11

RAPPLICATION NUMBER: 09/151889

RETLING DATE: 1998-09-11

RAPLICATION NUMBER: 09/169104

RETLING DATE: 1998-10-09

RETLING DATE: 1998-10-09
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R APPLICATION NUMBER: 08/828683
R FILING DATE: 1997-03-31
R APPLICATION NUMBER: 08/829270
R FILING DATE: 1997-03-31
R FILING DATE: 1997-09-11
R FILING DATE: 1997-09-11
                                      FILING DATE: 1999-02-12
APPLICATION NUMBER: 60/123972
FILING DATE: 1999-03-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 08/625328
FILING DATE: 1996-04-01
APPLICATION NUMBER: 08/710802
PILING DATE: 1996-09-23
APPLICATION NUMBER: 08/800699
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APPLICATION NUMBER: 09/304003
FILING DATE: 1999-04-30
APPLICATION NUMBER: 09/380137
FILING DATE: 1999-08-25
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FILING DATE: 1999-08-25
APPLICATION NUMBER: 09/380139
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APPLICATION NUMBER: 09/511133
FILING DATE: 2000-02-23
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APPLICATION UNMBER: 09/664610
FILING DATE: 2000-09-18
APPLICATION NUMBER: 09/665350
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    APPLICATION NUMBER: 60/119965
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FILING DATE: 1999-08-31
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APPLICATION NUMBER: 09/403297
FILING DATE: 1999-10-18
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APPLICATION NUMBER: 09/690169
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FILING DATE: 2000-11-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 08/934494
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APPLICATION NUMBER: 09/866034
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PRIOR
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TREATMENT OF TUNOR FILE REPERENCE: P2931ATA101

FILE REPERENCE: P2931ATA101

FILE REPERENCE: P2931ATA101

PRIOR PAPLICATION NUMBER: 60/01699

PRIOR FILING DATE: 1996-04-01

PRIOR FILING DATE: 1996-04-01

PRIOR FILING DATE: 1997-00-19

PRIOR FILING DATE: 1997-00-19

PRIOR FILING DATE: 1997-00-19

PRIOR FILING DATE: 1997-10-10

PRIOR FILING DATE: 1997-10-10

PRIOR FILING DATE: 1997-10-10

PRIOR FILING DATE: 1997-10-24

PRIOR PRILOR NUMBER: 60/066712

PRIOR PRILOR DATE: 1997-10-24

PRIOR PRILOR DATE: 1998-11-24

PRIOR PRILOR DATE: 1998-10-24

PRIOR PRILOR DATE: 1998-10-24

PRIOR PRILOR DATE: 1998-10-24

PRIOR PRILOR DATE: 1998-10-27

PRIOR PRILOR DATE: 1998-10-26

PRIOR FILING DATE: 1998-04-22

PRIOR PRILOR DATE: 1998-10-07

PRIOR PRILOR DATE: 1998-10-07
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                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
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APPLICATION NUMBER: 60/108867
FILING DATE: 1998-11-17
APPLICATION NUMBER: 60/112851
FILING DATE: 1998-12-16
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Smith, Victoria
Stone, Donna M.
Watanabe, Colin K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Godowski, Paul J.
Gurney, Austin L.
Hillan, Kenneth J
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Pan, James
Pitti, Robert M.
                                                                                                                                                                                                                                                                                   Query Match 69.3
Best Local Similarity 91.7
Matches 11; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wood, William I
                                                                             TYPE: DNA
CORGANISM: Homo sapiens
US-09-866-108-7717
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US-09-927-796-165
SEQ ID NO 7717
LENGTH: 17
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PRIOR APPLICATION NUMBER: 09/84733
PRIOR FILLING DATE: 2001-06-19
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PRIOR PELLING DATE: 1090-09-16
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PRIOR PELLING DATE: 1999-10-01
PRIOR PELLING DATE: 2000-02-18
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PRIOR APPLICATION NUMBER: PCT/US00/14941
PRIOR PELLING DATE: 2000-03-17
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PRIOR PELLING DATE: 2000-03-17
PRIOR PELLING DATE: 2000-03-17
PRIOR PELLING DATE: 2000-03-
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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/cgn2_6/ptodata/1/pna/US6028_COMB.seq:
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Copyright (c) 1993 - 2000 Compugen Ltd.
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Sequence 31, Appl Sequence 31, Appl Sequence 11, Appl Sequence 11, Appl Sequence 11, Appl Sequence 7, Appl Sequence 7, Appl Sequence 7, Appl Sequence 18071, Sequence 18071, Sequence 18070, Sequence 18070, Sequence 58705, Sequence 558705, Sequence 558705, Sequence 558705, Sequence 558705, Sequence 558705,

Description

Sequence 17, Appl. Sequence 17, Appl. Sequence 20465, A Sequence 417490, Sequence 61896, A Sequence 107363, Sequence 320910, Sequence 37434, Sequence 9034, Appl. Sequence 9034, Appl.

Sequence 22546, A Sequence 22546, A Sequence 22548, A Sequence 147526, Sequence 276612, Sequence 276137, Sequence 296787,

Sequence 365211, Sequence 536140, Sequence 536143,

Seguence 3, Appli Seguence 33931, A

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GENERAL INFORMATION:
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                                                                                                                                            APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Garade, William A.
APPLICANT: Garade, William A.
APPLICANT: Garade, William A.
APPLICANT: Nero, Pamela S.
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0476
CURRENT APPLICATION NUMBER: PCT/US00/18999
CURRENT APPLICATION NUMBER: PCT/US00/18999
CURRENT APPLICATION NUMBER: PCT/US00/18999
FRIOR PILLING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 31
LENGTH: 15
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GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
APPLICANT: Gened, William A.
TITLE OF INVENTION: Expression
TITLE OF INVENTION: Expression
FILE REPERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310A
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
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                ALIGNMENTS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CTHER INFORMATION: antisense sequence US-09-377-310-31
                                                                                                             Sequence 31, Application PC/TUS0018999 GENERAL INFORMATION:
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US-09-757-100B-31
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100.0%; Pr
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Matches 15; Conservative
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LENGTH: 15
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APPLICANT: Garde, William A.

APPLICANT: Garde, William A.

APPLICANT: Nero, Pamela S.

TITLE OF INVENTION: Expression

TITLE OF INVENTION: Expression

FILE REFERENCE: ISPH-0476

CURRENT APPLICATION NUMBER: PCT/US00/18999

CURRENT FILING DATE: 2000-07-13

PRIOR APPLICATION NUMBER: 09/377,310

PRIOR PRILING DATE: 1999-08-19

NUMBER OF SEQ ID NOS: 43

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 11

LENGTH: 20
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APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
APPLICANT: Nero, Pamela S.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase;
TITLE OF INVENTION: Expression
FILE REPREBLOE: ISPH-0533
CURRENT APPLICATION NUMBER: US/09/757,100B
CURRENT FILING DATE: 2001-03-15
PRIOR APPLICATION NUMBER: 09/377,310
PRIOR FILING DATE: 1999-08-19
PRIOR PILING DATE: 2000-07-13
NUMBER OF SEQ ID NOS: 44
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 31
LENGTH: 15
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100.0%; Pred. No. 2.5e+02;
Micmatches 0;
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; GENERAL INFORMATION:
; APPLICANT: ISIS Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                              ; OTHER INFORMATION: antisense sequence US-09-757-1008-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: antisense sequence PCT-US00-18999-11
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APPLICANT: Monia, Brett P.
                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 15; Conservative
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PCT-US00-18999-11
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Sequence 7, Application US/07918259
GENERAL INFORMATION:
APPLICANT: Hoke, Glenn D
APPLICANT: Bradley, Matthews O
APPLICANT: Williams, Taffy J
TITLE OF INVENTION: INFLAMMATORY COMPLICATIONS LIKE ARDS: ICAM-1 ANTISENSE
TITLE OF INVENTION: OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                             Gaps
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/918,259
FILIG DATE: 19920724
CLASSIFICATION: 514
                                                                                                                                                                                                                                                Score 13.4; DB 26;
Pred. No. 2.2e+03;
0; Mismatches 1;
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Pred. No. 8.3e+03;
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STREET: National Naval Medical Center
CITY: Bethesda
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                           NUMBER OF SEQ ID NOS: 140981
SOFWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 71111
LENGTH: 25
                                                                                                         , UMGANISM: HOMO Sapiens
; PUBLICATION INFORMATION:
; DATABASE ACCESSION NUMBER: GenBank S72904
US-09-660-220-71111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: N.C. 74,507
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 295-6759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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92.9%;
        PRIOR FILING DATE: 1999-11-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Spevack, A. David
REGISTRATION NUMBER: 24,7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (202) 295-4033
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                              Query Match 89.3
Best Local Similarity 93.3
Matches 14; Conservative
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Best Local Similarity 92.9
Matches 13; Conservative
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ZIP: 20889-5044
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy (
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US-07-918-259-7
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GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
APPLICANT: Garde, William A.
APPLICANT: Nero, Pamela S.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0533
CURRENT APPLICATION NUMBER: US/09/757,100B
CURRENT PILING DATE: 2001-03-15
PRIOR APPLICATION NUMBER: PCT/USO0/18999
PRIOR FILING DATE: 1999-08-19
PRIOR FILING DATE: 2000-07-13
NUMBER OF SEQ ID NOS: 44
SOFTWARE: PatentIn Ver. 2.0
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-03189
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEO ID NOS: 43
SOFTWARE: PATENTIN VET. 2.0
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GENERAL INFORMATION:
APPLICANT: Mittens et al.
APPLICANT: Affymetrix, inc.
TITLE OF INVENTION: Methods of Genetic Analysis of Human
FILE REFERENCE: 3102.1
CURRENT APPLICATION NUMBER: US/09/660,220
CURRENT FILIGE DATE: 2000-09-12
PRIOR APPLICATION NUMBER: 60/164,973
                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 15; DB 17; 100.0%; Pred. No. 2.5e+02;
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100.0%; Pred. No. 2.5e+02;
ive 0; Mismatches 0;
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; OTHER INFORMATION: antisense sequence
US-09-377-310-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CTHER INFORMATION: antisense sequence US-09-757-1008-11
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Best Local Similarity 100.
Matches 15; Conservative
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Best Local Similarity
Matches 15; Conserv
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                                                                                                                                                                SEQ ID NO 11
LENGTH: 20
                                                                                                                                                                                                          TYPE: DNA
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US-60-234-017-150700
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                                                           APPLICANT: ·Mittmann APPLICANT: ·Mittmann APPLICANT: Affymetrix, Inc. TITLE OF INVENTION: Methods of Genetic Analysis of the Rat TITLE OF INVENTION: Genome FILE REFERENCE: 3112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Affeweria, inc.
TITLE OF INVENTION: Methods of Genetic Analysis of the Rat
TITLE OF INVENTION: Genome
FILE REFERENCE: 3112
CURRENT APPLICATION NUMBER: US/60/233,166
CURRENT FILING DATE: 2000-10-24
NUMBER OF SEQ ID NOS: 420907
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 188071
LENGTH: 25
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GENERAL INFORMATION:
APPLICANT: Mittmann
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of the Rat
FITLE OF INVENTION: Genome
FILE REFERENCE: 3112
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92.9%; Pred. No. 8.5e+03;
Live 0; Mismatches 1;
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Pred. No. 8.5e+03;
0; Mismatches 1;
                                                                                                        CURRENT APPLICATION NUMBER: US/60/233,166
CURRENT APPLICATION NUMBER: US/60/233,166
CURRENT FILING DATE: 2000-10-24
NUMBER OF SEQ ID NOS: 420907
SOFTWARE: FastSEQ for Windows Version 4.0
EDO ID NO 72838
LENGTH: 25
                                                                                                                                                                                                                                                                                                 CRGANISM: Rattus norvegicus
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank AA819408
US-60-233-166-72838
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; DATABASE ACCESSION NUMBER: GenBank AA963963
US-60-233-166-188071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 188071, Application US/60233166
GENERAL INFORMATION:
APPLICANT: Mittmann
                    Sequence 72838, Application US/60233166 GENERAL INFORMATION:
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92.9%;
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Best Local Similarity 92.9
Matches 13; Conservative
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Matches 13; Conservative
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US-60-233-166-72838
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; Sequence 93096, Application US/60234017
; GENERAL INFORMATION:
; APPLICANT: Mittenan, M
; APPLICANT: Mittenan, M
; TITLE OF INVENTION: Methods of Genetic Analysis of Mus; TITLE OF INVENTION: Musculus
; TITLE PEFERRNCE: 3115
; CURRENT APPLICATION NUMBER: US/60/234,017
; CURRENT FILING DATE: 2000-09-20
; NUMBER OF SEQ ID NOS: 605887
; SOFTWARE: PASESEQ for Windows Version 4.0
; SEQ ID NO 93096
; LENGTH: 25
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Pred. No. 8.5e+03;
0; Mismatches 1;
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Pred. No. 8.5e+03;
0; Mismatches 1;
                                                                                                            ORGANISM: Rattus norvegicus
; PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank A1639536
US-60-233-166-293033
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; DATABASE ACCESSION NUMBER: GenBank AI854595
US-60-234-017-93096
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; DATABASE ACCESSION NUMBER: GenBank A1849180
US-60-234-017-150700
CURRENT APPLICATION NUMBER: US/60/233,166
CURRENT FILING DATE: 2000-10-24
NUMBER OF SEQ ID NOS: 420907
SEQ ID NO 293033
LENGTH: 25
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92.9%;
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Best Local Similarity 92.9
Matches 13; Conservative
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Best Local Similarity
Matches 13; Conserv
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Search completed: October 2, 2001, 21:50:21 Job time: 24534 sec

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Pred. No. 8.5e+03;
0; Mismatches 1; Indels (
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  Length 25;
                                                                                                                                                                                          US-60-234-017-150701
; Sequence 150701, Application US/60234017
; GENERAL INFORMATION:
; APPLICANT: Mitchann, M
; APPLICANT: Mitchann, M
; TITLE OF INVENTION: Methods of Genetic Analysis of Mus
; TITLE OF INVENTION: musculus
; FILE REFERENCE: 3115
; CURRENT APPLICANTON WUMBER: US/60/234,017
; CURRENT FILING DATE: 2000-09-20
; NUMBER OF SEQ ID NOS: 605887
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 150701
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 449268, Application US/60234017

Sequence 449268, Application US/60234017

GENERAL INFORMATION:
APPLICANT: Mittmann, M

TITLE OF INVENTION: Methods of Genetic Analysis of Mus
TITLE OF INVENTION: Methods of Genetic Analysis of Mus
TITLE REFERENCE: 3115
CURRENT APPLICATION NUMBER: US/60/234,017
CURRENT FILING DATE: 2000-09-20
NUMBER OF SEQ ID NOS: 605887
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 449268

LEMENTH: 25
                                            Indels
  Score 12.4; DB 55;
Pred. No. 8.5e+03;
0; Mismatches 1;
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PUBLICATION INFORMATION:
CATABASE ACCESSION NUMBER: Genbank AI849180
US-60-234-017-150701
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POBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank AW123353
US-60-234-017-449268
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92.9%;
Query Match 82.7%;
Best Local Similarity 92.9%;
Matches 13; Conservative
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Best Local Similarity 92.9
Matches 13; Conservative
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10 accaaggtgagtct 23
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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US-09-377-310-30
US-09-757-100B-30
PCT-US00-18999-10
US-09-377-310-10
US-09-757-100B-10
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US-09-263-959-1215
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/ Cgn2_6/ptodata/1/pna/US06_COMB.seq:*

/ Cgn2_6/ptodata/1/pna/US081_COMB.seq:*

/ Cgn2_6/ptodata/1/pna/US081_COMB.seq:*

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/ Cgn2_6/ptodata/1/pna/US082_COMB.seq:*

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/ Cgn2_6/ptodata/1/pna/US093_COMB.seq:*

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/1/pna/US6006_COMB.seq
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/cgn2_6/ptodata/1/pna/US6010_COMB.seq
/cgn2_6/ptodata/1/pna/US6011_COMB.seq
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     GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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/cgn2_6/
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                                                                                                                                                        October 2,
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Sequence 30, Appl

Description

Sequence 10, A Sequence 10, Sequence 30, Sequence 30,

Sequence 45, Appli Sequence 173, Appli Sequence 11530, Sequence 5, Appli Sequence 5, Appli Sequence 5, Appli Sequence 11530, Sequence 20692, A Sequence 20692, A Sequence 4119, Appli Sequence 4120, Appli Sequence 4120, Appli Sequence 4320, Appli Sequence 480, Appli Sequence 486, Appli

Sequence 486, App Sequence 487, App Sequence 241, App Sequence 7, Appli Sequence 1215, Ap Sequence 4, Appli Sequence 4, Appli Sequence 1215, Ap

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GENERAL INFORMATION:
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                                                                                                                                   APPLICANT: Isis Pharmaceuticals, Inc.
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Expression
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0476
CURRENT APPLICATION NUMBER: PCT/US00/18999
CURRENT FILING DATE: 2000-07-13
PRIOR FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: PatentIN Ver. 2.0
SEQ ID NO 30
LENGTH: 15
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GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
APPLICANT: Garde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310A
CURRENT FILING DATE: 1999-08-19
SOFTWARE: Patentin Ver. 2.0
SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
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100.0%; Pred. No. 4.5
+ive 0; Mismatches
                ALIGNMENTS
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                                                                                               Sequence 30, Application PC/TUS0018999 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 15; Conservative
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Best Local Similarity 100.
Matches 15; Conservative
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US-09-757-100B-30
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US-09-377-310-30
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Sequence 30, Application US/09757100B

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Monia, Prett P.
APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
APPLICANT: Nero, Pamela S.
TITLE OF INVENTION:
FILE REFERENCE: ISPH-0476
CURRENT APPLICATION NUMBER: PCT/US00/18999
CURRENT APPLICATION NUMBER: 09/377,310
PRIOR PILLING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 10
LENGTH: 20
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APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
APPLICANT: Nero, Pamela S.
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-053
CURRENT FILICATION NUMBER: US/09/757,100B
CURRENT FILING DATE: 2001-03-15
PRIOR APPLICATION NUMBER: 09/377,310
PRIOR APPLICATION NUMBER: PCT/US00/18999
PRIOR FILING DATE: 1999-0819
PRIOR FILING DATE: 2000-07-13
NUMBER OF SEQ ID NOS: 44
SOFTWARE: PATENTIN OF 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: antisense sequence US-09-757-100B-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Isis Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
COTHER INFORMATION: antisense sequence
PCT-US00-18999-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Sequence 10, Application PC/TUS0018999 ; GENERAL INFORMATION:
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; GENERAL INFORMATION:
; APPLICANT: Monia, Brett P.
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                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 15; Conservative
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Best Local Similarity 100.
Matches 15; Conservative
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PCT-US00-18999-10
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US-09-377-310-10
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Gaps
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GENERAL INFORMATION:
APPLICANT: The General Hospital Corporation
TITLE OF INVENTION: BETA NETRIN AND USES THEREOF
FILE REFERENCE: 10287-057W01
CURRENT APPLICATION NUMBER: PCT/US01/06413
CURRENT FILING DATE: 2001-02-28
PRIOR FILING DATE: 2000-02-29
PRIOR FILING DATE: 2000-02-29
PRIOR FILING DATE: 2000-01
SPRIOR FILING DATE: 2000-01
NUMBER OF SEQ ID NOS: 14
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 8
                                                                                                                                                                                                                                        SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 13.4; DB 7;
Pred. No. 3.7e+03;
0; Mismatches 1;
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                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/378,761
                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/987927
FILING DATE: 09-DEC-1992
ATTORNEY/AGENT IRFORMATION:
NAME: ZINDRICK, THOMAS D
                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                               ZINDRICK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CTHER INFORMATION: primer for PCR PCT-US01-06413-8
                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: ZINDRICK, THOMAS D
REGISTRATION NUMBER: 32185
REFERENCE/POCKET NUMBER: 38
TELECOMMUNICATION INFORMATION:
TELEPHONE: (517) 636-1869
INFORMATION FOR SEQ ID NO: 45:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Artificial Sequence
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93.3%;
                             E: THOMAS D. ZINI
9002 PURDUE ROAD
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Best Local Similarity 93.3
Matches 14; Conservative
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TYPE: nucleic acid
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      CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17 TAAGCAGCTGCAATT 3
                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
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                                        STREET: 9002 ...
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                                                                                                           COUNTRY: US
ZIP: 46268-1189
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                                                                                                                                                                                                                                                                                                         FILING DATE:
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PCT-US01-06413-8/c
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US-08-378-761-45
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APPLICANT: HEY, TIMOTHY D
APPLICANT: MORGAN, ALICE ER
TITLE OF INVENTION: RIBOSOME-INACTIVATING PROTEINS, INACTIVE
TITLE OF INVENTION: PRECURSOR FORMS THEREOF, A PROCESS FOR MAKING A METHOD OF
TITLE OF INVENTION: USING
NUMBER OF SEQUENCES: 59
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APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
TITLE OF INVENTION: Expression
TITLE OF INVENTION: 2001-03-15
CURRENT APPLICATION NUMBER: US/09/757,100B
CURRENT FILING DATE: 2001-03-15
PRIOR PPLICATION NUMBER: PST/US00/18999
PRIOR FILING DATE: 2000-07-13
NUMBER OF SEO ID NOS: 44
SOFTWARE: PatentIn Ver. 2.0
               TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase TITLE OF INVENTION: Expression FILE REFERENCE: 129H-0389
CURRENT APPLICATION NUMBER: US/09/377,310A
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 10
LENGTH: 20
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100.0%; Score 15; DB 29;
Best Local Similarity 100.0%; Pred. No. 4.8e+02;
Matches 15; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 15; DB 17; 100.0%; Pred. No. 4.8e+02;
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APPLICANT: Gaarde, William A.
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Best Local Similarity 100.
Matches 15; Conservative
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                                                                                                                                                                                                                    TYPE: DNA
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US-09-507-691-6/C

Sequence 6, Application US/09507691

TITER OF INVENTION: COLLAGEN BINDING PHYSIOLOGICALLY ACTIVE POLYPEPTIDE

TITER OF INVENTION: COLLAGEN BINDING PHYSIOLOGICALLY ACTIVE POLYPEPTIDE

CURRENT APPLICATION NUMBER: US/09/507,691

PRIOR FILING DATE: 1999-02-19

PRIOR FILING DATE: 1999-02-19

PRIOR FILING DATE: 1999-11-01

NUMBER OF SEQ ID NOS: 16

SOFTWARE: Patentin Ver. 2.0

TITER OF SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
OGANISM: Homo Sapiens
ORGANISM: Homo Sapiens
ORGANISM: Name/KEY: misc_feature
I CATTON: 1.27
US-09-338-907-173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 12.4; DB 17;
Pred. No. 1.4e+04;
0; Mismatches 1;
                                                      GENEKAL INFOGRATION:
APPLICANT: COCHEN, Daniel
APPLICANT: Blumenfeld, Marta
APPLICANT: Blumenfeld, Marta
APPLICANT: Blumenfeld, Marta
APPLICANT: Bougueleret, Lydie
TITLE OF INVENTION: PROSTATE CANCER GENE
FILE REFERENCE: GENSET.18CPLCP
CURRENT APPLICATION NUMBER: US/09/338,907
CURRENT APPLICATION NUMBER: 08/996,306
EARLIER FILING DATE: 1999-06-23
EARLIER FILING DATE: 1998-09-09
EARLIER FILING DATE: 1998-09-09
EARLIER FILING DATE: 1998-09-09
EARLIER FILING DATE: 1998-07-18
SOFTWARE: PALENT, DNOS: 578
SOFTWARE: PALENT, DNOS: 578
SEQ ID NO 173
                     Sequence 173, Application US/09338907
GENERAL INFORMATION:
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92.9%;
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Best Local Similarity 92.9
Matches 13; Conservative
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 25;
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                                                                                                     Sequence 76441. Application US/60234049
GENERAL INFORMATION:
APPLICANT: Mittmann, Michael
APPLICANT: Mittmann, Michael
APPLICANT: Affymetrix, Inc.;
TITLE OF INVENTION: Methods of Genetic Analysis of
TITLE OF INVENTION: Escherichia coli
FILE REFERENCE: 3117
CURRENT APPLICATION NUMBER: US/60/234,049
CURRENT FILING DATE: 2000-09-19
NUMBER OF SEQ ID NOS: 141629
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 76441
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 82.7%; Score 12.4; DB 55;
92.9%; Pred. No. 1.4e+04;
11ve. 0; Mismatches 1;
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GENERAL INFORMATION:
APPLICANT: Cohen, Daniel
APPLICANT: Cohen, Daniel
APPLICANT: Edumenfeld, Marta
APPLICANT: Blumenfeld, Marta
APPLICANT: Bougueleret, Lydie
TITLE OF INVENTION: Prostate cancer gene
FILE REFERENCE: GENSET. 018CP1
CURRENT FILING DATE: 1997-12-22
EARLIER APPLICATION NUMBER: 08/996,306
EARLIER FILING DATE: 1997-12-22
EARLIER FILING DATE: 1998-09-09
NUMBER OF SEQ ID NOS: 578
SOFTWARE: PATENT. PM
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank U00096
1; 660-234-049-76441
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Best Local Similarity 92.9%;
Matches 13; Conservative,
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LOCATION: 1..27
21 AAGCAGCTGCCCTT
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Best Local Similarity
Matches 13; Conserva
                                                                                       US-60-234-049-76441/C
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US-09-218-207-173/c
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LENGTH: 27
                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
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Length 27;

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FEATURE:
COTHER INFORMATION: Description of Artificial Sequence: PCR sense
COTHER INFORMATION: primer for human fibronectin collagen-binding
COTHER INFORMATION: domain
COT-09-507-691-6
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                                                                                                                                                 Length 49;
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                                                                                                                                             82.7%; Score 12.4; DB 19;
92.9%; Pred. No. 1.6e+04;
tive 0; Mismatches 1;
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Length 25; Indels

80.0%; Score 12; DB 55; I 100.0%; Pred. No. 2.4e+04; tive 0; Mismatches 0;

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TITLE OF INVENTION: Methods of Genetic Analysis of TITLE OF INVENTION: Arabidopsis thaliana
                                                                                                                                                                                TYPE: DNA
CRGANISM: Arabidopsis thaliana
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank AL035394
US-60-233-620-111530
                                    FILE REFERENCE: 3116
CURRENT APPLICATION NUMBER: US/60/233,620
CURRENT FILING DATE: 2000-10-24
NUMBER OF SEQ ID NOS: 131820
SOFTWARE: FastSEQ for Windows Version 4.0
LENGTH: 25
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Best Local Similarity
Matches 12; Conserv
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Pred. No. 2.2e+04;
1; Mismatches 1; Indels
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APPLICANT: Mittmann
APPLICANT: Affymetrix, Inc.; TITLE OF INVENTION: Methods of Genetic Analysis of; TITLE OF INVENTION: Arabidopsis thaliana; FILE REFERENCE 3116
CURRENT APPLICATION NUMBER: US/60/233,620; CURRENT FILING DATE: 2000-10-24; NUMBER OF SEQ ID NOS: 131820; SOGTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                         APPLICANT: Chew, Anne APPLICANT: Chew, Anne APPLICANT: Choi, Julie APPLICANT: Choi, Julie APPLICANT: Choi, Julie APPLICANT: Choi, Julie APPLICANT: Genaissance Pharmaceuticals, Inc. TITLE OF INVENTION: Haplotypes of the PONI Gene FILE REFERENCE: PONIPCT CURRENT APPLICATION NUMBER: PCT/USO1/05126 CURRENT FILING DATE: 2001-02-16 PRIOR FILING DATE: 2000-02-16 PRIOR FILING DATE: 2000-02-16 SPIOR FILING DATE: 2000-02-16 SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               80.0%; Score 12; DB 55; I larity 100.0%; Pred. No. 2.4e+04; Conservative 0; Mismatches 0;
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; DATABASE ACCESSION NUMBER: Genbank AC007591
US-60-233-620-95891
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APPLICANT: Mittmann
APPLICANT: Affymetrix, Inc.
                                                                            Sequence 13, Application PC/TUS0105126 GENERAL INPORMATION:
PPELICANT: Denton, Richard R
APPLICANT: Nandabalan, Krishnan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Arabidopsis thaliana
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85.7%;
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Best Local Similarity 85.7
Matches 12; Conservative
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Best Local Similarity
Matches 12; Conserv
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US-60-233-620-111530
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US-60-233-620-95891
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LENGTH: 15
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Sequence 10084, A
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1. /cgn2_6/ptodata/2/pna/PCT_NEW_COMB.seq:*
2. /cgn2_6/ptodata/2/pna/USO6_NEW_COMB.seq:*
3: /cgn2_6/ptodata/2/pna/USO7_NEW_COMB.seq:*
4: /cgn2_6/ptodata/2/pna/USO8_NEW_COMB.seq:*
5: /cgn2_6/ptodata/2/pna/USO9_NEW_COMB.seq:*
6: /cgn2_6/ptodata/2/pna/USO9_NEW_COMB.seq:*
7: /cgn2_6/ptodata/2/pna/USO9_NEW_COMB.seq2:*
8: /cgn2_6/ptodata/2/pna/USO9_NEW_COMB.seq2:*
GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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S-09-866-108-7019
S-09-866-108-7019
S-09-866-108-7021
S-09-866-108-11910
S-09-866-108-11911
S-09-866-108-11911
S-09-866-108-11912
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S-09-866-108-11918
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US-09-708-690-10084
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US-09-923-515-15
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                                                                                                                         2, 2001, 16:55:06
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Maximum Match 100%
Listing first 45 summaries
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Match Length
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26 11.4 76.0 20 5 US-09-923-515-18 Sequence 18, Appl 28 11.4 76.0 20 6 US-09-198-452A-2269 Sequence 2269, Ap 29 11.4 76.0 20 6 US-09-198-452A-4764 Sequence 4764, Ap 29 11.4 76.0 47 5 US-09-198-772-315-7 Sequence 10, Appl 29 11.4 76.0 47 5 US-09-772-315-7 Sequence 10, Appl 20 23 11 73.3 17 6 US-09-866-108-7016 Sequence 7016, Ap 20 23 11 73.3 20 1 PCT-US01-18763A-187 Sequence 7016, Appl 20 23 11 73.3 20 1 PCT-US01-18763A-189 Sequence 119.08, Appl 20 23 11 73.3 20 1 PCT-US01-18763A-189 Sequence 119.08, Appl 20 2 2 US-09-866-108-119.08 Sequence 12, Appl 20 2 2 US-09-866-108-119.08 Sequence 119.08, Appl 20 2 US-09-866-108-119.08 Sequence 119.08, Appl 20 2 US-09-866-108-119.08 Sequence 119.08, Appl 20 2 US-09-866-108-119.08 Sequence 20 2 Appl 20 2 US-09-866-108-119.08 Sequence 20 2 Appl 20 2 US-09-766-095-78 Se
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18:90-86-108-7017 Application US/09866108
Sequence 7017 Application US/09866108
Sequence 7017 Application US/09866108
SPEZAL INFORMATION:
APPLICARY: 01, Yongang
APPLICARY: BANNEL, David R.
APPLICARY: BANNEL, David R.
APPLICARY: BANNEL, David R.
APPLICARY: SHANNON, MARK.
TITLE OF INVENTION: MOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
CURRENT APPLICATION WIMBER: US/09/466.108
CURRENT APPLICATION MUMBER: US/09/466.108
CURRENT APPLICATION WIMBER: US/09/466.108
FRIOR FILING DATE: 2000-05-26 60/207,456
PRIOR PLING DATE: 2000-00-26 60/207,456
PRIOR PLING DATE: 2000-00-26 60/207,456
PRIOR PLING DATE: 2000-00-27 60/205
PRIOR PLING DATE: 2000-00-27 60/205
PRIOR PLING DATE: 2000-00-30 60/205
PRIOR PLING DATE: 2000-00-30 60/205
PRIOR PLING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/0066
PRIOR PLING DATE: 2001-01-30
PRIOR PLING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/0066
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR PRIOR PRIOR DATE: 2001-01-30
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APPLICANT: SHANNOW, MARK

TITLE REPERENCE: ABOMICA-7

TITLE REPERENCE: ABOMICA-7

CURRENT APPLICATION NUMBER: US/09/866,108

CURRENT FILING DATE: 2001-05-25

PRIOR FILING DATE: 2001-05-26

PRIOR FILING DATE: 2000-10-30

PRIOR FILING DATE: 2000-10-30

PRIOR FILING DATE: 2000-10-30

PRIOR FILING DATE: 2001-01-30

PRIOR PRIOR FILING DATE: 2001-01-30

PRIOR PRILOR PRILOR NUMBER: PCT/USO1/00661

PRIOR PRILOR DATE: 2001-01-30

PRIOR PRILOR DATE: 2001-01-30

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PRIOR PRILOR DATE: 2001-01-30

PRIOR PRILOR PRILOR DATE: 2001-01-30

PRIOR PRILOR PRILOR NUMBER: PCT/USO1/00670

PRIOR PRILOR PRILOR DATE: 2001-01-30

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Pred. No. 2.7e+03;
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10.09-86-108-7020/c

1. Sequence 7020, Application US/09866108

2. GENERAL INFORMATION:

3. APPLICANT: GU, Yizhong
                                                                                                                                                                                                                                             Sequence 7019, Application US/09866108 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                        APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
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Best Local Similarity 100.
Matches 12; Conservative
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; ORGANISM: Homo sapiens
US-09-866-108-7019
          2 aagcagctgcca 13
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US-09-866-108-7019/c
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SOFTWARE: ACOUL
SEQ ID NO 7019
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APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
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CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2001-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-10-10-10
PRIOR FILING DATE: 2000-10-10-10
PRIOR FILING DATE: 2000-10-10-10
PRIOR FILING DATE: 2001-01-30
                                                                                                                                                                                                      80.0%; Score 12; 100.0%; Pred. No.
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US-09-866-108-7018/C
Sequence 7018, Application US/09866108
GENERAL INFORMATION:
APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
                                                                                                                                                                                                   Query Match
Best Local Similarity 100.0
Matches 12; Conservative
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; LENGTH: 17
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-7017
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; ORGANISM: Homo sapiens
US-09-866-108-7018
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Best Local Similarity
Matches 12; Conserva
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APPLICANT: GU, Yizhong APPLICANT: JI, Yonggang APPLICANT: GU, Yizhong APPLICANT: JI, Yonggang APPLICANT: BENN, Sharron G. APPLICANT: PENN, Sharron G. APPLICANT: HANZEL, David R. APPLICANT: RANK, David R. APPLICANT: CHEN, Wensheng APPLICANT: SHANNON, MAIK TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE FILE REFERENCE: AZONICA-7 CURRENT APPLICATION NUMBER: US 60/207, 456

PRIOR APPLICATION NUMBER: US 60/207, 456

PRIOR APPLICATION NUMBER: GB 24263.6

PRIOR APPLICATION NUMBER: QS 60/207, 456

PRIOR APPLICATION NUMBER: PUS 60/236, 359

PRIOR APPLICATION NUMBER: PUS 60/236, 359

PRIOR PILING DATE: 2000-09-27

PRIOR APPLICATION NUMBER: POT/US01/00666

PRIOR APPLICATION NUMBER: POT/US01/00667

PRIOR APPLICATION NUMBER: POT/US01/00667
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PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR PELLING DATE: 2000-10-04
PRIOR PELLING DATE: 2000-10-03
PRIOR PELLING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00667
PRIOR PELLING DATE: 2001-01-30
PRIOR PELLING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00669
PRIOR PELLING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00669
PRIOR PELLING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Acomica Sequence Listing Engine
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Matches 12; Conservative
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US-09-866-108-7021
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US-09-866-108-7022/c
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                                                                                                                                  ### PRPLICANT; CHEN, Wenshing
### PRILICANT; CHEN, Wenshing
### PRILICANT; CHEN, Wenshing
#### PRILICANT; CHEN, Wenshing
#### CURRENT APPLICANTON; WYCSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCIE
#### CURRENT APPLICANTON UNDBER; US/09/866,108
#### CURRENT APPLICANTON UNDBER; US/09/866,108
#### PRILING DATE: 2000-10-26
### PRILING DATE: 2000-10-34
### PRILING DATE: 2000-10-34
### PRILING DATE: 2000-10-30
### PRILING DATE: 2000-10-30
### PRILING DATE: 2001-01-30
### PRILING
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APPLICANT: JI, Yonggang
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANEL, David K.
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: WYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
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SOFTWARE: Acomica Sequence Listing Engine
SEQ ID NO 7020
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       PENN, Sharron G.
HANZEL, David K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 80.0
Best Local Similarity 100.
Matches 12; Conservative
                                                                                   RANK, David R. CHEN, Wensheng
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CORGANISM: Homo sapiens
US-09-866-108-7020
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APPLICANT: CHANNON, MEARY
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
TITLE REFERENCE: ADOMICA-7
CURRENT APPLICATION UNMER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
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. 2.9e+03; ·
ches 0;
                 PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2000-09-21
PRIOR PILING DATE: 2000-09-21
PRIOR PILING DATE: 2000-09-21
PRIOR PILING DATE: 2001-09-25
NUMBER OF SEQ ID NOS: 15752
SEQ ID NOS: 15752
SEQ ID NO 11909
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CURRENT FILING DATE: 2001-05-25
PRIOR PILING DATE: 2000-05-26
PRIOR PELING DATE: 2000-05-26
PRIOR PELING DATE: 2000-10-04
PRIOR PELING DATE: 2000-10-07
PRIOR PELING DATE: 2000-09-27
PRIOR PILING DATE: 2001-01-30
PRIOR PELING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00665
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; Sequence 11910, Application US/09866108
; GENERAL INFORMATION:
; APPLICANT: GU, Yizhong
; APPLICANT: JI, Yonggang
; APPLICANT: HANZEL, David K.
; APPLICANT: RANK, David R.
; APPLICANT: CHEN, Wensheng
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PRIOR APPLICATION NUMBER: PCT/USO1/00663
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00662
PRIOR FLING DATE: 2001-01-30
PRIOR PLICATION DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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APPLICATION NUMBER: US 60/266,860
FILING DATE: 2001-02-05
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Best Local Similarity 100.0%;
Matches 12; Conservative C
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; ORGANISM: Homo sapiens
US-09-866-108-11909
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GREEKAL INFORMATION:

APPLICANT: GU, YIZHONG

APPLICANT: GU, YIZHONG

APPLICANT: GU, YIZHONG

APPLICANT: TI, YONGGANG

APPLICANT: PENN, Sharron G.

APPLICANT: RANK, David R.

APPLICANT: RANK, David R.

APPLICANT: RANK, David R.

APPLICANT: RANK, Wensheng

APPLICANT: CHEN, Wensheng

APPLICANT: SALNON, MAIK

TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE

FILE REPERENCE: BODMICA-7

CURRENT FILING DATE: 2001-05-25

FRIOR FILING DATE: 2000-05-26

FRIOR PLICATION NUMBER: US 60/207,456

FRIOR PELICATION NUMBER: CB 24263.6

FRIOR PELICATION NUMBER: PCT/US01/0066

FRIOR APPLICATION NUMBER: PCT/US01/0066

FRIOR PLICATION NUMBER: PCT/US01/0066

FRIOR PLICATION NUMBER: PCT/US01/0066

FRIOR PLICATION NUMBER: PCT/US01/0066

FRIOR FILING DATE: 2001-01-30

FRIOR FILING DATE: 2001-01-30
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100.0%; Pred. No. 2.7e+03;
tive 0; Mismatches 0;
                 PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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PRIOR PILING DATE: 2001-01-30
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PRIOR PILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR PILING DATE: 2000-09-21
PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Acomica Sequence Listing Engine
SEG ID NO 7022
APPLICATION NUMBER: PCT/US01/00664
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Best Local Similarity 100.
Matches 12; Conservative
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; ORGANISM: Homo sapiens
US-09-866-108-7022
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GENERAL INFORMATION:
APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David R.
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, MAIK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
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100.0%; Pred. No. 2.9e+03;
.ive 0; Mismatches 0;
       O; Mismatches 0;
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LENGTH: 25
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       100.0%;
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Best Local Similarity 100.
Matches 12; Conservative
                                         12; Conservative
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                                                                                                          2 aagcagctgcca 13
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       Best Local Similarity
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APPLICANT: D1, YONGGANION;
APPLICANT: PUN, Sharron G.
APPLICANT: HANEEL, David K.
APPLICANT: HANEEL, David K.
APPLICANT: HANEEL, David R.
APPLICANT: CHEN, Wensheng
FILE REPRENCE: APPLICANTON, MAIR
TITLE OF INVENTION: MYOSIE: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR PELICATION NUMBER: US 60/207,456
PRIOR PLICATION NUMBER: US 60/236,359
PRIOR PELICATION NUMBER: US 60/236,359
PRIOR PLICATION NUMBER: D3 60/207,466
PRIOR FLING DATE: 2000-09-27
PRIOR PLICATION NUMBER: PCT/USO1/00666
PRIOR FLILING DATE: 2001-01-30
PRIOR PLICATION NUMBER: PCT/USO1/00666
PRIOR PLICATION NUMBER: PCT/USO1/00667
PRIOR PLICATION NUMBER: PCT/USO1/00669
PRIOR PLICATION NUMBER: PCT/USO1/00660
PRIOR PLICATION NUMBER: PCT/USO1/00660
PRIOR PLICATION NUMBER: PCT/USO1/00660
PRIOR PLICATION NUMBER: US 60/234,687
PRIOR PLICATION NUMBER: US 60/234,687
PRIOR PLICATION NUMBER: US 60/26,860
PRIOR PRIOR APPLICATION NUMBER: US 60/26,860
PRIOR PRIOR APPLICATION NUMBER: US 60/26,860
PRIOR PRIOR APPLICATION NUMBER: US 60/200,800
PRIOR PLICATION NUMBER: US 60/2000
PRIOR PLICATION NUMBER: US
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Pred. No. 2.9e+03;
NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Aeomica Sequence Listing Engine
SEQ ID NO 11910
LENGTH: 25
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Best Local Similarity 100.
Matches 12; Conservative
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ORGANISM: Homo sapiens
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RESULT 11 US-09-866-108-11913/c

Length 25;

80.0%; Score 12; DB 6;

Query Match

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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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LENGTH: 25
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GENERAL INFORMATION:
APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANK, DAVIG K.
APPLICANT: RANK, David K.
APPLICANT: RANK, David R.
Sequence 11913, Application US/09866108 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity 100
Matches 12; Conservative
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US-09-866-108-11913
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US-09-866-108-11914/C
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APPLICANT: GU, YIZHONG
APPLICANT: JI, Yonggang
APPLICANT: BENN, Barron G.
APPLICANT: HANZEL, David R.
APPLICANT: HANZEL, David R.
APPLICANT: RANK, David R.
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE ADOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT PILING DATE: 2001-05-26
PRIOR PRILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
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PRIOR APPLICATION NUMBER: US 60/236,359
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CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT PELING DATE: 2001-05-26
PRIOR PELING DATE: 2000-05-26
PRIOR PELING DATE: 2000-05-26
PRIOR PELING DATE: 2000-10-04
PRIOR PELING DATE: 2000-10-04
PRIOR PELING DATE: 2000-10-27
PRIOR PELING DATE: 2000-10-30
PRIOR FILING DATE: 2001-01-30
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SOFTWARE: Aeomica Sequence Listing Engine
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; ORGANISM: Homo sapiens
US-09-866-108-11914
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US-09-866-108-11915/c
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GENERAL INFORMATION:

APPLICANT: GU, Yizhong
APPLICANT: H. Yongquag
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT APPLICATION NUMBER: US/0207,456
PRIOR PILING DATE: 2001-05-25
PRIOR PAPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-010-04
PRIOR FILING DATE: 2000-010-04
PRIOR FILING DATE: 2000-010-04
PRIOR PILING DATE: 2001-01-30
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PRIOR PILING DATE: 2001-01-30
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PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-31
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PRIOR FILING DATE: 2001-01-31
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PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
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PPLICATION NUMBER: PCT/US01/00670
FILING DATE: 2001-01-30
APPLICATION NUMBER: US 60/234,687
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Best Local Similarity 100.
Matches 12; Conservative
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; ORGANISM: Homo sapiens
US-09-866-108-11916
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US-09-866-108-11917/c
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GENERAL INFORMATION:
APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: RANK, David R.
APPLICANT: SHANNON, WENSHONG
APPLICANT: SHANNON, WARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILLE REPERENCE: AEGMICA-7
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CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-03-36,359
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2001-01-30
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; ORGANISM: Homo sapiens
US-09-866-108-11915
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Sequence 229423, Sequence 229555, Sequence 279728, Sequence 279728, Sequence 279728, Sequence 157, App Sequence 157, App Sequence 157, App Sequence 235736, Sequence 235736, Sequence 273976, Seq
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Sequence 3, Appli
Sequence 3, Appli
Sequence 37375, A
Sequence 37376, A
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Sequence 23, Appl
Sequence 23, Appl
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Sequence 68034
Sequence 13699
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Sequence 7906,
Sequence 14258
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US-60-233-166-279728
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US-60-234-017-42928

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US-60-234-017-273976

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US-60-234-017-376806

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US-09-006-298-11
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Copyright (c) 1993 - 2000 Compugen Ltd.
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                                                                                Sequence 23, Application PC/TUS0018999
GENERAL INFORMATION:
APPLICANT: Isis Pharmaceuticals, Inc.
APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Moro, Pamela S.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase;
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0476
CURRENT APPLICATION NUMBER: PCT/US00/18999
CURRENT FILING DATE: 2000-07-13
PRIOR FILING DATE: 1999-08-19
PRIOR FILING DATE: 1999-08-19
NUMBER: OF SEQ ID NOS: 43
SOFTWARE: PatentIN Ver. 2.0
SEQ ID NO 23
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APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT FILIG APPLICATION NUMBER: US/09/377,310A
CURRENT FILIG DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SECTIANE: Patentin Ver. 2.0
SEQ ID NO 23
LENGTH: 15
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llarity 100.0%; Pred. No. 3.4e+02;
Conservative 0; Mismatches 0;
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100.0%; Pred. No. 3.4e+02;
tive 0; Mismatches 0;
ALIGNMENTS
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Best Local Similarity 100.
Matches 15; Conservative
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US-09-757-100B-23
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#### APPLICANT: Monia, Brett P.
### APPLICANT: Monia, Brett P.
#### APPLICANT: Monia, Brett P.
#### APPLICANT: More, Pamela S.
#### TITLE OF INVENTION: Expression
### TITLE OF INVENTION: DATE: 2000-07-13
### PRIOR PAPLICATION NUMBER: 09/377,310
### PRIOR APPLICATION NUMBER: 09/377,310
### PRIOR PAPLICATION NUMBER: 09/377,310
### SEQ ID NOS: 43
### SEQ ID NOS: 43
### SEQ ID NO 3
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APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Monia, Milliam A.
APPLICANT: Mero, Pamela S.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0533
CURRENT APPLICATION NUMBER: US/09/757,100B
FILE REFERENCE: 2001-03-15
PRIOR FILING DATE: 1999-08-19
PRIOR APPLICATION NUMBER: OP/377,310
PRIOR FILING DATE: 1999-08-19
PRIOR FILING DATE: 1999-08-19
PRIOR FILING DATE: 2000-07-13
NUMBER OF SEO ID NOS: 44
SOFTWARE: PatentIn Ver. 2.0
SEO ID NO 23
LENGTH: 15
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Pred. No. 3.4e+02;
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APPLICANT: Isis Pharmaceuticals, Inc.
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US-09-377-310-3
'Sequence 3, Application US/09377310A
'GENERAL INFORMATION:
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Best Local Similarity 100.
Matches 15; Conservative
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GENERL INFORMATION:
APPLICANT: Mitthann et al.
TITLE OF INVENTION: Methods of Genetic Analysis of Human; FILE REFERENCE: 3102.1
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GENERAL INFORMATION:
APPLICANT: Mittment al.
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of Human
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Pred. No. 1.1e+04;
); Mismatches 1;
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Pred. No. 1:1e+04;
0; Mismatches 1
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NUMBER OF SEQ ID NOS: 140981
SOFTWARE: FastSEQ for Windows Version 4.0
                             SEQ ID NOS: 140981
FastSEQ for Windows Version 4.0
                                                                                                                                                    ; PUBLICATION INFORMATION:
; DATABASE ACCESSION NUMBER: GenBank L11931
US-09-660-220-37375
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; DATABASE ACCESSION NUMBER: GenBank L11931
US-09-660-220-37376
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DATABASE ACCESSION NUMBER: Genbank L11931
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            1999-11-11
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Best Local Similarity 92.9
Matches 13; Conservative
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                                                                                                              TYPE: DNA
ORGANISM: Homo Sapiens
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ORGANISM: Homo Sapiens
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ORGANISM: Homo Sapiens
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          PRIOR FILING DATE: 1'
NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ fo
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Best Local Similarity
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US-09-660-220-37376
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LENGTH: 25
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APPLICANT: Gaarde, William A.
APPLICANT: Garde, William A.
APPLICANT: Mero, Pamela S.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
TITLE OF INVENTION: Expression
TITLE OF INVENTION: Expression
TITLE OF INVENTION: 2001-03-15
FILE REFERENCE: ISPH-053
CURRENT APPLICATION NUMBER: US/09/757,100B
PRIOR APPLICATION NUMBER: 091-03-15
PRIOR PELLING DATE: 1999-048-19
PRIOR FILING DATE: 2000-07-13
NUMBER OF SEQ ID NOS: 44
SOFTWARE: PatentIn Ver. 2.0
APPLICANT: Gaarde, William A.

TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310A
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 3
LENGTH: 20
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APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of Human
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CURRENT APPLICATION NUMBER: US/09/660,220
CURRENT FILING DATE: 2000-09-12
PRIOR APPLICATION NUMBER: 60/164,973
                                                                                                                                                                                                                                                                        OTHER INFORMATION: antisense sequence US-09-377-310-3
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Best Local Similarity 100.0%;
Matches 15; Conservative (
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                                                                                                                                                                                                                 TYPE: DNA
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Gaps

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US-60-233-166-296555/c
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LENGTH: 25
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                                                                                                                                                                                                                                                        TYPE: DNA
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APPLICANT: Mitthmann

APPLICANT: Affymetrix, Inc.

TITLE OF INVENTION: Methods of Genetic Analysis of the Rat

TITLE OF INVENTION: Genome

FILE REFERENCE: 3112

CURRENT APPLICATION NUMBER: US/60/233,166

CURRENT APPLICATION NUMBER: US/60/233,166

NUMBER OF SEQ ID NOS: 420907

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 229423

LENGTH: 25
                                                   Length 25;
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                                                                                                                                                                                                                              Sequence 136549, Application US/60232638
GENERAL INFORMATION:
APPLICANT: Mittmann
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of Yeast
FILE REFREENCE: 3110
CURRENT APPLICATION NUMBER: US/60/232,638
CURRENT FILING DATE: 2000-09-14
NUMBER OF SEQ ID NOS: 138410
SOUTHWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 136549
LENGTH: 25
                                                                                  Indels
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                                                Score 12.4; DB 26;
Pred. No. 1.1e+04;
0; Mismatches 1;
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Pred. No. 1.1e+04;
0; Mismatches 1;
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Pred. No. 1.1e+04;
0; Mismatches 1;
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; DATABASE ACCESSION NUMBER: GenBank AF016901
US-60-233-166-229423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; PUBLICATION INFORMATION:
; DATABASE ACCESSION NUMBER: SGD YPR119W
US-60-232-638-136549
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Saccharomyces Cerevisea
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                                                  82.7%;
92.9%;
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92.9%;
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                                              Query Match 82.7
Best Local Similarity 92.9
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 82.7
Best Local Similarity 92.9
Matches 13; Conservative
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Best Local Similarity
Matches 13; Conserv
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US-60-233-166-229423
US-09-660-220-37377
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Sequence 296555, Application US/60233166
GENERAL INFORMATION:
APPLICANT: Mitumann
APPLICANT: Mitumann
APPLICANT: Mitumann
TITLE OF INVENTION: Methods of Genetic Analysis of the Rat
TITLE OF INVENTION: Genome
FILE REFERENCE: 3112
CURRENT APPLICATION NUMBER: US/60/233,166
CURRENT APPLICATION NUMBER: US/60/233,166
CURRENT FILING DATE: 2000-10-24
NUMBER OF SEQ ID NOS: 420907
SEQ ID NO 296555
LENGTH: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 25;
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Mittmann
APPLICANT: Mittmann
APPLICANT: Mittmann
APPLICANT: Mittmann
APPLICANT: Mittmann
APPLICANT: Mittmann
APPLICANT: Methods of Genetic Analysis of Rat
TITLE OF INVENTION: Methods of Genetic Analysis of Rat
FILE REFERENCE: 3114
CURRENT APPLICATION NUMBER: US/60/233,357
CURRENT FILING DATE: 2000-10-24
NUMBER OF SEQ ID NOS: 21305
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 297601, Application US/60234017
GENERAL INFORMATION:
APPLICANT: Mittmann, M
APPLICANT: Affiguetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of Mus
TITLE OF INVENTION: musculus
FILE REFERENCE: 3115
CURRENT APPLICATION NUMBER: US/60/234,017
CURRENT FILING DATE: 2000-09-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 12.4; DB 55;
Pred. No. 1.1e+04;
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Pred. No. 1.1e+04;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                        ORGANISM: Rattus norvegicus
PUBLICATION INFORMATION:
JATABASE ACCESSION NUMBER: Genbank A1639536
US-60-233-166-296555
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; DATABASE ACCESSION NUMBER: Genbank AF016901
US-60-233-357-1753
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  82.78;
92.98;
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92.9%;
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Matches 13; Conservative
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Best Local Similarity
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Search completed: October 2, 2001, 21:50:20 Job time: 24533 sec

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5161, 5162, 5163, 5164,

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TITLE OF INVENTION: ANTISENSE MODULATION OF RhoG EXPRESSION
FILE REFERENCE: FYSP-0106
CURRENT APPLICATION NUMBER: US/09/787,252
CURRENT FILING DATE: 2001-03-15
PRIOR APPLICATION NUMBER: 09/161,015
PRIOR FILING DATE: 1998-09-25
NUMBER OF SEQ ID NOS: 47
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
             US-09-735-271-1792
US-09-245-025-21
US-09-245-025-21
US-09-245-025-21
US-09-708-690-20444
US-09-866-108-229
US-09-866-108-229
US-09-866-108-223
US-09-866-108-223
US-09-866-108-213
US-09-866-108-213
US-09-866-108-5158
US-09-866-108-5158
US-09-866-108-5160
US-09-866-108-5160
US-09-866-108-5160
US-09-866-108-5160
US-09-866-108-5160
US-09-866-108-5160
US-09-866-108-5160
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                                                                                                                                                                                                                                                                  US-09-866-108-5161
US-09-866-108-5162
US-09-866-108-5163
US-09-866-108-5164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 2037, Application US/09735271
GENERAL INFORMATION:
APPLICANT: Daly, Mark J.
APPLICANT: Hudson, Thomas J.
APPLICANT: Lander, Eric S.
APPLICANT: Rioux, John
APPLICANT: Siminovitch, Kathy
TITLE OF INVENTION: IBD-RELATED POLYMORPHISMS
FILE REFERENCE: 2825.1025-002
CURRENT APPLICATION NUMBER: US/09/735,271
CURRENT FILING DATE: 2000-12-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: Antisense Oligonucleotide US-09-787-252-17
                                                                                                                                                                                                                                                                                                                                                                      ALIGNMENTS
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PRIOR FILING DATE: 1999-12-10
PRIOR APPLICATION NUMBER: US 60/196,046
PRIOR FILING DATE: 2000-04-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 17, Application US/09787252
; GENERAL INFORMATION:
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                                                    Best Local Similarity 92.3
Matches 12; Conservative
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   US-09-787-252-17/c
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Query Match
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                                                                                                                                    (without alignments)
14.467 Million cell updates/sec
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15629,
15630,
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15635,
15636,
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Sequence 15633,
                                                                                                                 ; Search time 2173.58 Seconds
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Sequence
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1: /cgn2_6/ptodata/2/pna/PCT_NEW_COMB.seq:*

2: /cgn2_6/ptodata/2/pna/USOG_NEW_COMB.seq:*

3: /cgn2_6/ptodata/2/pna/USOB_NEW_COMB.seq:*

5: /cgn2_6/ptodata/2/pna/USOB_NEW_COMB.seq:*

6: /cgn2_6/ptodata/2/pna/USOB_NEW_COMB.seq:*

7: /cgn2_6/ptodata/2/pna/USOB_NEW_COMB.seq:*

7: /cgn2_6/ptodata/2/pna/USOB_NEW_COMB.seq:*

8: /cgn2_6/ptodata/2/pna/USOB_NEW_COMB.seq2:*

8: /cgn2_6/ptodata/2/pna/USOB_NEW_COMB.seq2:*
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Compugen Ltd.
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US-09-866-108-10732

US-09-866-108-10733

US-09-866-108-10734

US-09-866-108-10735

US-09-866-108-10736

US-09-925-548-93

US-09-925-548-93

US-09-866-108-15624

US-09-866-108-15626

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US-09-866-108-15634
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US-09-866-108-15636
                                                                                                                                                                                                                                                                                                    1895189 seqs, 1048201267 residues
                                                                                                                                                                                                                                                                                                                                   Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SUMMARIES
             GenCore version Copyright (c) 1993 - 2000
                                                                                                                 October 2, 2001, 16:55:06
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Maximum Match 100%
Listing first 45 summaries
                                                                                  - nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
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15
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Match Length
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Sequence:
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SEQ ID NO 10732
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APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
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                                                                                                                                                            Score 11.4; DB 7; Length 25;
Pred. No. 5.3e+03;
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Pred. No. 5.5e+03;
0; Mismatches 1; Indels
                                                                                                                                                                                                   Indels
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GENERAL INFORMATION:
APPLICANT: Dander, Thomas J.
APPLICANT: Lander, Eric S.
APPLICANT: Rioux, John
APPLICANT: Rioux, John
APPLICANT: Siminovitch, Kathy
TITLE OF INVENTION: IBD-RELATED POLYMORPHISMS
FILE REPERENCE: 2825.1025-002
                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/735,271
CURRENT FILING DATE: 2000-12-11
FRIOR APPLICATION NUMBER: US 60/170,257
PRIOR FILING DATE: 1999-12-10
PRIOR FILING DATE: 2000-04-10
NUMBER OF SEQ ID NOS: 2058
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1558
LENGTH: 40
SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 2037
LENGTH: 25
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GENERAL INFORMATION:
APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, DAVIG K.
APPLICANT: RANK, David K.
APPLICANT: CHEN, Wensheng
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PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
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; OTHER INFORMATION: n = A,T,C or G
US-09-735-271-1558
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92.3%;
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92.3%;
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Best Local Similarity 92.3
Matches 12; Conservative
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Best Local Similarity 92.3
Matches 12; Conservative
                                                    TYPE: DNA
COGANISM: Homo sapiens
US-09-735-271-2037
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US-09-866-108-10732
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APPLICANT: RANK. David R.
APPLICANT: CHEN, Wensheng
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, MAIK
ITILE ON INVENTION: WOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US 60/207,456
PRIOR PILING DATE: 2000-105-25
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
PRIOR PRIOR PAPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
PRIOR PILICATION NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
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Pred. No. 2e+04;
); Mismatches
PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-10-37
PRIOR FILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR PLING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00661
PRIOR PLING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR APPLICATION NUMBER: DCT/US01/00670
PRIOR APPLICATION NUMBER: US 60/256,860
PRIOR PLING DATE: 2000-09-21
PRIOR PLING DATE: 2000-09-21
PRIOR PLING DATE: 2000-09-21
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ilarity 91.7%;
Conservative
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US-09-866-108-10732
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6 gggctcccagtg 17
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Best Local Similarity
Matches 11; Conserv
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APPLICANT: RANK, David R.
APPLICANT: RANK, David R.
APPLICANT: RANK, Wensheng
APPLICANT: SHANNON, MARK
TITLE OF INVENTION: WINDSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR PELICATION NUMBER: GB 24263.6
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2001-01-30
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Pred. No. 2e+04;
n. Mismatches 1; Indels
                      PRIOR APPLICATION NUMBER: PCT/US01/00670
PRIOR FLLING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR PELLOR DATE: 2000-09-21
PRIOR PELLOR DATE: 2000-09-21
PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Aeomica Sequence Listing Engine
SEQ ID NO 10734
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R APPLICATION NUMBER: PCT/USO1/00669

R APPLICATION NUMBER: PCT/USO1/00665

R FILING DATE: 2001-01-30

R APPLICATION NUMBER: PCT/USO1/00663

R APPLICATION NUMBER: PCT/USO1/00663

R APPLICATION NUMBER: PCT/USO1/00663
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SOFTWARE: Acomica Sequence Listing Engine
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APPLICATION NUMBER: PCT/US01/00661
ELING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00670
FILING DATE: 2001-01-30
    APPLICATION NUMBER: PCT/US01/00661
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APPLICATION NUMBER: US 60/266,860
FILING DATE: 2001-02-05
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Best Local Similarity 91.7
Matches 11; Conservative
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CORGANISM: Homo sapiens
US-09-866-108-10734
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Pred. No. 2e+04;
0; Mismatches 1;
             PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00665
PRIOR APPLICATION NUMBER: PCT/USO1/00668
PRIOR APPLICATION NUMBER: PCT/USO1/00668
PRIOR PILING DATE: 2001-01-30
PRIOR PRILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR PRILING DATE: 2001-01-30
PRIOR PLING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR PLING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00670
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR APPLICATION NUMBER: US 60/266,860
PRIOR PRILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
APPLICATION NUMBER: PCT/US01/00669
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PRIOR APPLICATION NUMBER: PCT/US01/00662
PRIOR FILING DATE: 2001-01-30
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Sequence 10734, Application US/09866108
GENERAL INFORMATION:
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91.7%;
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PENN, Sharron G.
HANZEL, David K.
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Best Local Similarity 91.7
Matches 11; Conservative
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; ORGANISM: Homo sapiens
US-09-866-108-10733
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APPLICANT: CHEN, Marsh ABOUT IN WEASHERS
APPLICANT: CHEN, Warsh ABOUT ATON NUMBER: PCT/USO1/00663

PRIOR FILING DATE: 2001-01-30

PRIOR RELIEVED ABOUT ABOUT
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Pred. No. 2e+04;
); Mismatches
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; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
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Best Local Similarity 91.7
Matches 11; Conservative
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CHEN, Wensheng
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ORGANISM: Homo sapiens
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US-09-925-548-93/c
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
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Pred. No. 2e+04;
); Mismatches 1;
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR PLING DATE: 2000-10-04
PRIOR PILING DATE: 2000-10-04
PRIOR PILING DATE: 2000-10-130
PRIOR PPLICATION NUMBER: PCT/USO1/00667
PRIOR PELICATION NUMBER: PCT/USO1/00667
PRIOR PILING DATE: 2001-01-30
PRIOR PELING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 10736, Application US/09866108
GENERAL INFORMATION:
APPLICANT: GU, Yizhong
APPLICANT: JT, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANSEL, DBVIG K.
APPLICANT: RANK, David K.
APPLICANT: CHEN, Wensheng
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LENGTH: 17
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PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
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Best Local Similarity 91.7%;
Matches 11; Conservative
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; SEQ ID NO 10735
; LENGTH: 17
: TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-10735
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; ORGANISM: Homo sapiens
US-09-866-108-10736
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Best Local Similarity
Matches 11; Conserv
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US-09-866-108-10736
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Gaps

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APPLICANT: HARK, David R.

APPLICANT: HARK, David R.

APPLICANT: GENEW, Wensheng
APPLICANT: GENEW, Wensheng
APPLICANT: GIRNON, MARK

FILE REPERROKE: ABOMICA-6

CURREMY PILING DATE: 2001-05-25

CURREMY PELLORION NUMBER: US 60/207,456

PRIOR FILING DATE: 2000-05-26

PRIOR PLING DATE: 2000-09-27

PRIOR PLING DATE: 2000-09-27

PRIOR PLING DATE: 2000-09-27

PRIOR PLING DATE: 2000-09-27

PRIOR PPLICATION NUMBER: PCT/USO1/0066

PRIOR PPLICATION NUMBER: PCT/USO1/0066

PRIOR APPLICATION NUMBER: PCT/USO1/0066

PRIOR PLING DATE: 2001-01-30

PRIOR PLIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 10.4; DB 6; Length 21;
Pred. No. 2e+04;
1; Mismatches 2; Indels
                  APPLICANT: LIGHTER, CETC S.

TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
FILE REFERENCE: 2825.2008-001
CURRENT APPLICATION NUMBER: US 60/176,861
PRIOR APPLICATION NUMBER: US 60/176,861
PRIOR FILING DATE: 2000-01-19
NUMBER OF SEQ ID NOS: 461
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 111
LENGTH: 21
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SOFTWARE: Acomica Sequence Listing Engine
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APPLICATION NUMBER: US 60/234,687
FILING DATE: 2000-09-21
APPLICATION NUMBER: US 60/266,860
FILING DATE: 2001-02-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 15624, Application US/09866108
; GENERAL INFORMATION:
; APPLICANT: GU, Yizhong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  69.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 cgggctcacagtgg 15
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; ORGANISM: Homo sapiens
US-09-765-081-111
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Best Local Similarity
Matches 11; Conserv
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Pred. No. 2e+04;
0; Mismatches 1; Indels
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      APPLICANT: Decudat, SHOLDANT: Pecudat, Greg
APPLICANT: Yee, Arthur
TYTE OF INVENTION: INTEGRIN-LINKED KINASE AND ITS USES
FILE REFERENCE: KINEOOLCIP4
CURRENT APPLICATION NUMBER: US/09/925,548
CURRENT FILING DATE: 1999-09-03
PRIOR APPLICATION NUMBER: 09/390,425
PRIOR FILING DATE: 1999-09-03
PRIOR APPLICATION NUMBER: 09/035,706
PRIOR FILING DATE: 1998-03-05
PRIOR APPLICATION NUMBER: 08/955,841
PRIOR FILING DATE: 1996-11-19
PRIOR FILING DATE: 1996-11-19
PRIOR FILING DATE: 1996-11-19
PRIOR PLILING DATE: 1996-11-19
PRIOR APPLICATION NUMBER: 60/009,074
PRIOR FILING DATE: 1995-11-19
SOFTWARE: FASTSEQ for Windows Version 4.0
SOFTWARE: FASTSEQ for Windows Version 4.0
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APPLICANT: Macdonald Morris
APPLICANT: Macdonald Morris
APPLICANT: Tom Ryder
APPLICANT: Tom Ryder
APPLICANT: Tom Ryder
APPLICANT: Town Ryder
APPLICANT: Town Ryder
APPLICANT: Town Ryder
TILE OF INVENTION: Town Nucleic Acids and Probe Arrays
FILE REFERENCE: 3108
CURRENT APPLICATION NUMBER: US/09/827,383
CURRENT APPLICATION NUMBER: US 60/195,585
PRIOR FILING DATE: 2000-04-06
NUMBER OF SEQ ID NOS: 2050
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1980
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 10.4; DB 5;
Pred. No. 2e+04;
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Best Local Similarity 91.7%;
Matches 11; Conservative
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Best Local Similarity 91.7
Matches 11; Conservative
APPLICANT: Dedhar, Shoukat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-925-548-93
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SPEIGNERAL INFORMATION:

APPLICANT: GIV, Yizhong
APPLICANT: TIN, Yongsang
APPLICANT: TIN, Yongsang
APPLICANT: TIN, Yongsang
APPLICANT: THANZEL, David K.
APPLICANT: THANZEL, David K.
APPLICANT: THANZEL, David K.
APPLICANT: CHEN, Mensheng
APPLICANT: CHEN, Mensheng
APPLICANT: CHEN, Mensheng
APPLICANT: GINVENTION: MYONGRE: US 60204.456
FILE REBERREER APPLICANTON NUMBER: US 60204.456
FILE REBERREER APPLICANTON NUMBER: US 60204.456
FILE REPERREER APPLICANTON NUMBER: US 60204.456
FILE APPLICANTON NUMBER: PCT/US01/00666
FILE APPLICANTON NUMBER: PCT/US01/00666
FILE APPLICANTON NUMBER: PCT/US01/00669
FILE APPLICANTON NUMBER: PCT/US01/00669
FILE APPLICANTON NUMBER: PCT/US01/00669
FILE APPLICANTON NUMBER: PCT/US01/00669
FILE APPLICANTON NUMBER: PCT/US01/00661
FILE APPLICANTON NUMBER: PCT/US01/00670
FILE APPLICANTON NUMBER: US 60/246,860
FILE APPLICANTON NUMBER: US 60/246,860
FILE APPLICANTER APPLICANTER APPLICANTER SEQUENCE LIST
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Pred. No. 2e+04;
0; Mismatches 1;
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                                                                                                                                                                                                                                                            Sequence 15626, Application US/09866108 GENERAL INFORMATION:
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91.7%;
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Best Local Similarity 91.7
Matches 11; Conservative
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US-09-866-108-15626
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US-09-866-108-15626
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APPLICANT: 01, Yizhong
APPLICANT: 01, Vizhong
APPLICANTION WHERE: 02, 01, 01, 02, 02, 0466, 108
CURRENT FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-06-26
PRIOR FILING DATE: 2000-06-36
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/0665
PRIOR APPLICATION NUMBER: PCT/US01/0667
PRIOR APPLICATION NUMBER: PCT/US01/0670
PRIOR APPLICATION NUMBER: PC
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Pred. No. 2e+04;
0; Mismatches 1; Indels
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                                                                                                                                                                                                                        69.3%;
91.7%;
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Best Local Similarity 91.7
Matches 11; Conservative
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                                                           ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-866-108-15624
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; ORGANISM: Homo sapiens
US-09-866-108-15625
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14 gggctcccagtg 25
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Best Local Similarity
Matches 11; Conserv
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SEQ ID NO 15624
LENGTH: 25
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16.701 Million cell updates/sec
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(cgn2_6/ptodata1/pna/US084_COMB.seq:*

(cgn2_6/ptodata1/pna/US084_COMB.seq:*

(cgn2_6/ptodata1/pna/US086_COMB.seq:*

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(cgn2_6/ptodata1/pna/US087_COMB.seq:*

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(cgn2_6/ptodata1/pna/US095_COMB.seq:*

(cgn2_6/ptodata1/pna/US095_COMB.seq:*
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/cgn2_6/prodate/1/pna/US096C_COMB.Seq:
/cgn2_6/prodate/1/pna/US096C_COMB.Seq:
/cgn2_6/prodate/1/pna/US096C_COMB.Seq:
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/cgn2_6/prodate/1/pna/US000_COMB.Seq:
/cgn2_6/prodate/1/pna/US000_COMB.Seq:
/cgn2_6/prodate/1/pna/US000_COMB.Seq:
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/cgn2_6/ptodata/1/pna/US6011_COMB.seq:*
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/ptodata/1/pna/US6007_COMB.seq:
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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14155048 segs, 7388405095 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                               - nucleic search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 catgatgcttaaaagcttac 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IDENTITY_NUC
Gapop 10.0 , Gapext 1.0
                                                                                                                                                                                                                                                                                                                                                                                          US-09-757-100B-20
20
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Maximum DB seq length: 50
                                                                                                                                                                                                                                                                                                                                                                                                   Title:
Perfect score:
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/cgn2_6/ptodata/1/pna/US6020_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6020_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6022_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6023_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6024_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6025_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6025_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6025_COMB.seq:*
                 /cgn2_6/ptodata/1/pna/US6013_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6015_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6015_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6016_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6016_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6016_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6019_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6019_COMB.seq:*
                                                                                                                                                                                                                                                                                                                  /cgn2_6/ptodata/1/pna/US6027_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6028_COMB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

r i	duence	<u>`</u>	quence 40,	e 40,	equence 40,	equence 498317,	equence 8, 1	equence 443	Sequence 123801,	equence 564	equence 564	564	137	13/		00	82143,	Š	7	4	9	Sequence 5612, Ap	, TZ99	Sequence 6632, Ap	82138,	m	42140	4214	equence 430178,	o a	140	14,	21,	e 21,	21	equence 25,	equence 21,	ednence 38,	23,	Sequence 24, Appl
SUMMARIES	T-US00-18999-2	US-03-377-310-20	JS00-18999-40	-09-377-310-4	09-757-100B-40	50-234-017-	09-261-301C-8	50-233-166-4432	0-233-620-1238 0-234-012-5645	50-234-017-5645	50-234-017-5645	50-234-017-5645	50-232-638-1373	50-237-038-13/3 50-234-017-2040	38-818-604A-5	09-345-828-8	50-232-638-	3-224-593-5	09-509-668-4	50-234-690-4034	50-232-638-684	0-232-638-661	50-232-638-662	50-232-638-6632	50-233-620-821	50-233-620-8214	50-234-017-4214	50-234	50-234-017-4301	30-234-049-4694 3-584-040-329	5-09-465-684-1	08-484-174-14	r-us98-01468-2	S-09-013-896-2	09-013-896A-	s-09-216-599-2	-827-948-	S-09-216-599-3	-09-216-599-2	US-09-216-599-24
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GENERAL INFORMATION:
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                                                                                                                                       APPLICANT: Isls Pharmaceuticals, Inc.
APPLICANT: State Pharmaceuticals, Inc.
APPLICANT: Garde, William A.
APPLICANT: Garde, William A.
APPLICANT: Garde, William A.
APPLICANT: Nero, Pamela S.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase TITLE OF INVENTION: Expression
FILE REPERENCE: ISPH-0476
CURRENT FILING DATE: 2000-07-13
PRIOR FILING DATE: 1999-08-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REPERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310A
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SEQ ID NO 20
LENGTH: 20
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Pred. No. 11;
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                 ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 20;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION: antisense sequence PCT-US00-18999-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CTHER INFORMATION: antisense sequence US-09-377-310-20
                                                                                                      Sequence 20, Application PC/TUS0018999 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-377-310-20; Sequence 20, Application US/09377310A; GENERAL INFORMATION:
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US-09-757-100B-20
; Sequence 20, Application US/09757100B
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Best Local Similarity 100.0%;
Matches 20; Conservative 0;
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 43
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 20
LENGTH: 20
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Best Local Similarity 100.
Matches 20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Monia, Brett P.

APPLICANT: Gaarde, William A.

APPLICANT: Bardel B.

TITLE OF INVENTION: Expression

TITLE OF INVENTION: Expression

FILE REFRENCE: ISPH-0476

CURRENT APPLICATION WUMBER: PCT/US00/18999

CURRENT APPLICATION WUMBER: PCT/US00/18999

PRIOR APPLICATION WUMBER: 09/377,310

PRIOR PILING DATE: 1999-08-19

NUMBER OF SEQ ID NOS: 43

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0533
CURRENT APPLICATION NUMBER: 09/377,100B
CURRENT APPLICATION NUMBER: 09/377,310
PRIOR APPLICATION NUMBER: PCT/US00/18999
PRIOR FILING DATE: 1999-08-19
PRIOR FILING DATE: 2000-07-13
NUMBER OF SEQ ID NOS: 44
SCOFFWARE: Patentin Ver. 2.0
SEQ ID NO 20
LENGTH: 20
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'n. 38+03;
0; Indels
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: antisense sequence US-09-757-100B-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Isis Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: antisense sequence PCT-US00-18999-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 40, Application PC/TUS0018999 ; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Sequence 40, Application US/09377310A; GENERAL INFORMATION:
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100.0%; Pre
0;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 100.0%; Some Best Local Similarity 100.0%; P. Matches 20; Conservative 0;
                                                                                                                                                                                                                                                                                                           TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 catgatgcttaaaagcttac 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 15; Conservative
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PCT-US00-18999-40
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US-09-377-310-40
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Gaps

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Indels

Length 25;

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APPLICANT: Abbort Laboratories
APPLICANT: Abbort Laboratories
APPLICANT: Abbort Laboratories
APPLICANT: Abbort Laboratories
APPLICANT: Cohen, Gerald
APPLICANT: Cohen, Gerald
APPLICANT: *Usiel, Orit
APPLICANT: *Usiel, Orit
TITLE OF INVENTION: Thereof Useful As Antimicrobial Agents
TITLE OF INVENTION: Thereof Useful As Antimicrobial Agents
FILE REFERENCE: 6286.US.01
CURRENT APPLICATION NUMBER: US,09/261,301C
CURRENT APPLICATION NUMBER: US 60/076,525
PRIOR FILING DATE: 1999-03-02
NUMBER OF SEQ ID NOS: 10
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Mittmann
APPLICANT: Mittmann
APPLICANT: Affymerrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of the Rat
TITLE OF INVENTION: Genome
FILE REFERENCE: 3112
CURRENT FILING DATE: 2000-10-24
NUMBER OF SEQ ID NOS: 420907
SECTIONE FASTSEQ for Windows Version 4.0
SEQ ID NO 44325
LENGTH: 25
                                                                                                                                                                                                                                                                        71.0%; Score 14.2; DB 55;
84.2%; Pred. No. 8e+03;
11ve 0; Mismatches 3;
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88.2%; Pred. No. 1.3e+04;
tive 0; Mismatches 2;
                                                                                                                                                             ; PUBLICATION INFORMATION:
; DATABASE ACCESSION NUMBER: GenBank AV213434
US-60-234-017-498317
NUMBER OF SEQ ID NOS: 605887
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 498317
LENGTH: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-60-233-166-44325; Sequence 44325, Application US/60233166; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: C-terminal primer US-09-261-301C-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 8, Application US/09261301C GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                            7 atactgcttaaaagtttac 25
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Best Local Similarity 84.2
Matches 16; Conservative
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 8
US-09-261-301C-8/C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
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APPLICANT: Gaarde, William A.

APPLICANT: Gaarde, William A.

APPLICANT: Gaarde, William A.

PAPLICANT: Monia, Brett P.

TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase;

TITLE OF INVENTION: Expression

FILE PERERNCE: ISPH-0533

CURRENT APPLICATION NUMBER: 05/09/757,100B

CURRENT FILING DATE: 2001-03-15

PRIOR FILING DATE: 1999-08-19

PRIOR FILING DATE: 2000-07-13

NUMBER OF SEQ ID NOS: 44

SOFFWARE: Patentin Ver. 2.0

SEQ ID NO 40
                     APPLICANT: Gaarde, William A.

TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310A
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
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GENERAL INFORMATION:
A APPLICANT: Mittmann, M
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of Mus
TITLE OF INVENTION: musculus
FILE REFERENCE: 3115
CURRENT APPLICATION NUMBER: US/60/234,017
CURRENT FILING DATE: 2000-09-20
                                                                                                                                                                                                                                                                                                                                                                                                       Score 15; DB 17;
; Pred. No. 3e+03;
0; Mismatches 0
                                                                                                                                                                                                                                                                                         FEATURE:
; OTHER INFORMATION: antisense sequence
US-09-377-310-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: antisense sequence US-09-757-100B-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 40, Application US/09757100B GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 75.0%;
Best Local Similarity 100.0%;
Matches 15; Conservative 0
                                                                                                                                                                                                                                                 TYPE: DNA ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 15; Conservative
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Indels

Length 27;

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US-60-234-017-564527/c; Sequence 564527, Application US/60234017; GENERAL INFORMATION:
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   1 catgatgcttaaaagcttac 20
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80.0%;
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80.0%;
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                      Mittmann, M
Affymetrix, Inc.
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Best Local Similarity 80.09
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Best Local Similarity
Matches 16; Conserv
                                                                                      RESULT 12
US-60-234-017-564521/C
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APPLICANT:
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GENERAL INFORMATION:
APPLICANT: Mittmann, M
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of Mus
TITLE OF INVENTION: Musculus
FILE REFERENCE: 3115
CURRENT APPLICATION NUMBER: US/60/234,017
CURRENT FILING DATE: 2000-09-20
NUMBER OF SEQ ID NOS: 605887
SEQ ID NO 564512
LENGTH: 25
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Sequence 123801, Application US/60233620

SEDWERL INFORMATION:

APPLICANT: Mittmann

APPLICANT: Affymetrix, Inc.

TITLE OF INVENTION: Methods of Genetic Analysis of

TITLE OF INVENTION: Arabidopsis thaliana

FILE REFERENCE: 3116

CURRENT APPLICATION NUMBER: US/60/233,620

CURRENT FILIG DATE: 2000-10-24

NUMBER OF SEQ ID NOS: 131820

SEQ ID NO 123801

LENGTH: 25
                                                                                                                       Query Match

68.0%; Score 13.6; DB 55;
Best Local Similarity 80.0%; Pred. No. 1.6e+04;
Matches 16; Conservative 0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            68.0%; Score 13.6; DB 55;
80.0%; Pred. No. 1.6e+04;
11ve 0; Mismatches 4;
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80.0%; Pred. No. 1.6e+04;
Live 0; Mismatches 4;
                                 ; PUBLICATION INFORMATION:
; DATABASE ACCESSION NUMBER: GenBank AA818914
US-60-233-166-44325
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; DATABASE ACCESSION NUMBER: Genbank AV245764
US-60-234-017-564512
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PUBLICATION INFORMATION:
CS-0-233-620-123801
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TYPE: DNA ORGANISM: Rattus norvegicus
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Best Local Similarity 80.0°
Matches 16; Conservative
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Best Local Similarity
Matches 16; Conserv
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Sequence 564521, Application US/60234017
GENERAL INFORMATION:
APPLICANT: Mitthann, M
TITLE OF INVENTION: Methods of Genetic Analysis of Mus
TITLE OF INVENTION: Methods of Cenetic Analysis of Mus
TITLE OF INVENTION: Methods of Cenetic Analysis of Mus
FILE REFERENCE: 3115
CURRENT APPLICATION NUMBER: US/60/234,017
CURRENT FILING DATE: 2000-09-20
NUMBER OF SEQ ID NOS: 605887
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 564521
LENGTH: 25
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US-60-234-017-564524/C
Sequence 564524, Application US/60234017
GENERAL INFORMATION:
APPLICANT: Wittmann, M
APPLICANT: Wittmann, M
TITLE OF INVENTION: Methods of Genetic Analysis of Mus
TITLE OF INVENTION: Methods of Cenetic Analysis of Mus
FILE REPERBENCE: 3115
CURRENT APPLICATION WUMBER: US/60/234,017
CURRENT FILING DATE: 2000-09-20
NUMBER OF SEQ ID NOS: 605887
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 565524
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Pred. No. 1.6e+04;
0; Mismatches 4;
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Pred. No. 1.6e+04;
0; Mismatches 4;
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; DATABASE ACCESSION NUMBER: GenBank AV245764
US-60-234-017-564521
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; DATABASE ACCESSION NUMBER: GenBank AV245764
US-60-234-017-564524
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US-60-232-638-137320/c
; Sequence 137320, Application US/60232638
; GENERAL INFORMATION:
; APPLICANT: Mittmann
APPLICANT: Mittmann
TITLE OF INVENTION: Methods of Genetic Analysis of Yeast
FILE REFERENCE: 3110
CURRENT APPLICATION NUMBER: US/60/232,638
; CURRENT FILING DATE: 2000-09-14
; NUMBER OF SEQ ID NOS: 138410
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 137320
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 93.3%; Pred. No. 2e+04;
Matches 14; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                               Indels
TITLE OF INVENTION: Methods of Genetic Analysis of MusrITLE OF INVENTION: musculus FILE REFERENCE: 3115
CURRENT APPLICATION NUMBER: US/60/234,017
CURRENT FILING DATE: 2000-09-20
NUMBER OF SEQ ID NOS: 605887
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 564527
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                                                                                                                                                                       TYPE: DNA
CRGANISM: Mus musculus
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank AV245764
US-60-234-017-564527
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SOCHATOMYCES CETEVISEA
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: SGD YPR165W
US-60-232-638-137320
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COUNTRY: USA
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US-09-536-784-418/c
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; ORGANISM: HOMU
US-09-715-849-159
LENGTH: 21
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19.290 Million cell updates/sec
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                                                                                               Search time 2173.58 Seconds
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2. /cgn2_6/ptodata/2/pna/USOG_NEW_COMB.seq:*
3: /cgn2_6/ptodata/2/pna/USOR_NEW_COMB.seq:*
4: /cgn2_6/ptodata/2/pna/USOB_NEW_COMB.seq:*
5: /cgn2_6/ptodata/2/pna/USOB_NEW_COMB.seq:*
6: /cgn2_6/ptodata/2/pna/USOB_NEW_COMB.seq:*
7: /cgn2_6/ptodata/2/pna/USOB_NEW_COMB.seq:*
8: /cgn2_6/ptodata/2/pna/USOB_NEW_COMB.seq2:*
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US-09-536-784-418
US-09-765-272-318
US-09-765-272-330
US-60-253-778-315-44
US-60-295-722-60-6
US-60-295-722-60-6
US-60-295-722-60-6
US-09-537-784-272
US-09-765-272-272
US-09-765-272-316
US-09-591-306-33
US-09-591-306-33
US-09-310-8448-328
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US-09-310-8448-346
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US-60-253-456-6918
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             GenCore version
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GENERAL INFORMATION:
APPLICANT: Choi et. al.
TITLE OF INVENTION: Streptococcus pneumoniae Antigens and Vaccines NUMBER OF SEQUENCES: 452
CORRESPONDENCE ADDRESS:
ADDRESSE: Human Genome Sciences, Inc.
STREET: 9410 Key West Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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Sequence 3
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                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Cargill, Michele
APPLICANT: Ireland, James S.
APPLICANT: Lander. Erelc S.
TTTLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
FILE REFERENCE: 2825.2002-001
CURRENT APPLICATION NUMBER: US/09/715,849
CURRENT PAPLICATION NUMBER: US/000-11.17
PRIOR APPLICATION NUMBER: US 60/167,334
NUMBER OF SEQ ID NOS: 589
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 20850
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage
COMPUTER: HP Vectra 486/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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Pred. No. 4.3e+03;
; Mismatches 3;
        US-09-546-745A-1852
US-09-546-745A-1853
US-09-920-759-70
US-09-920-759-70
US-09-9375-318-66
US-09-465-589-6
US-09-465-589-6
US-09-935-370-118
US-09-935-370-118
US-09-938-370-118
US-09-801-274-1514
US-09-801-274-1514
US-09-465-589-15
US-09-664-6108-345
US-09-804-65-89-15
US-09-804-661-4
US-09-804-661-4
US-09-804-61-4
US-09-804-61-4
                                                                                                                                                                                                                                                                                           ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                        ; Sequence 159, Application US/09715849; GENERAL INFORMATION:
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77.8%;
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Matches 14; Conservative
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Sequence 330, Application US/09536784
GENERAL INFORMATION:
APPLICANT: Choi et. al.
TITLE OF INVENTION: Streptococcus pneumoniae Antigens and Vaccines NUMBER OF SEQUENCES: 452
CORRESPONDENCE ADDRESS:
ADDRESSE: Human Genome Sciences, Inc.
STREET: 9410 Key West Avenue
CITY: Rockville
STATE: Maryland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 330, Application US/09765272
GENERAL INFORMATION:
APPLICANT: Choi et. al.
TITLE OF INVENTION: Streptococcus pneumoniae Antigens and Vaccines NUMBER OF SEQUENCES: 452
                                                                                                                                                           Gaps
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                                                                                                                   Length 36;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
COMPUTER REDEE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage
COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS Version 6.2
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
                                                                                                                                                         Indels
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                                                                                                              Score 12.6; DB 6;
Pred. No. 5.8e+03;
); Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 12.2; DB 5;
Pred. No. 9.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/09/536,784
FILING DATE: 30-Oct-1997
CLASSIFICATION: <UNKNOWN>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 08/961,083
FILING DATE: OCT-30-1997
ATTORNEY/AGENT INFORMATION:
NAME: Michelle S. Marks
REGISTRATION NUMBER: 41,971
REFERENCE/DOCKET NUMBER: PB340P3
TELECOMMUNICATION INFORMATION:
                  ; SEQUENCE DESCRIPTION: SEQ ID NO: 418: US-09-765-272-418
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE DESCRIPTION: SEQ ID NO: 330: US-09-536-784-330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (301) 309-8504
TELEFAX: (301) 309-8512
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 33 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 330 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    61.0%;
82.4%;
                                                                                                              Query Match 63.0%;
Best Local Similarity 78.9%;
Matches 15; Conservative
                                                                                                                                                                                                    2 atgatgcttaaaagcttac 20
                                                                                                                                                                                                                                               21 ATTATGTTGATAAGCTTAC 3
      TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4 gatgcttaaaagcttac 20
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Best Local Similarity
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US-09-765-272-330/c
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; Sequence 418, Application US/09765272
; Sequence 418, Application US/09765272
; GENERAL INFORMATION:
; APPLICANT: Choi et. al.
; TITLE OF INVENTION: Streptococcus pneumoniae Antigens and Vaccines NUMBER OF SEQUENCES: 452
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage COMPUTER: HP Vectra 486/33 OPERATING SYSTEM: MSDOS version 6.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 12.6; DB 5;
Pred. No. 5.8e+03;
0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sciences, Inc.
OPERATING SYSTEM: MSDOS version 6.2
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
                                                                 APPLICATION NUMBER: US/09/536,784 FILING DATE: 30-0ct-1997 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/09/765,272
FILING DATE: 22-Jan-2001
CLASSIFICATION: <UNKNOWN>
                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: PB340P3
TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8514
INFORMATION FOR SEQ ID NO: 418:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Brookes, A. Anders
REGISTRATION NUMBER: 36,373
REFERENCE/DOCKET NUMBER: PB340P2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE DESCRIPTION: SEQ ID NO: 418: US-09-536-784-418
                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/961,083
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 08/961,083
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Human Genome Scien
STREET: 9410 Key West Avenue
                                                                                                                                                                                                                  NAME: Michelle S. Marks
REGISTRATION NUMBER: 41,971
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                              FILING DATE: OCT-30-1997
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                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 36 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 36 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: ASCII TEXT CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 418
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 63.0%;
Best Local Similarity 78.9%;
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS
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COMPUTER READABLE FORM:
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GEMERAL INFORMATION.

APPLICANT: Barbara Bour
APPLICANT: Blake Denison
APPLICANT: Blake Denison
APPLICANT: Brancar Bihain
APPLICANT: Bernard Bihain
APPLICANT: Lydie Bougueleret
TITLE OF INVENTION: Thereof
FILE REFERENCE: 104.US2.PRO
CURRENT APPLICATION NUMBER: US/60/295,600
CURRENT APPLICATION DATE: 2001-06-04
PRIOR FILING DATE: 2001-06-04
NUMBER OF SEQ ID NOS: 18
SOFTWARE: Patent.pm
SEQ ID NO 6
                                                                                          APPLICANT: JOHN Chicca
APPLICANT: Blake Denison
APPLICANT: Blake Denison
APPLICANT: Frances Yen
APPLICANT: Frances Yen
APPLICANT: Jydie Bougueleret
TITLE OF INVENTION: GSSP4 Polynucleotides and Polypeptides and Uses Thereof
FILE REFERENCE: 103.US2.PRO
CURRENT APPLICATION NUMBER: US/60/295,722
CURRENT FILING DATE: 2001-06-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 12.2; DB 8;
Pred. No. 9.6e+03;
); Mismatches 3;
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Pred. No. 9.6e+03;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 6, Application US/60295600 ; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 6, Application US/60295722; GENERAL INFORMATION:
APPLICANT: Barbara Bour
APPLICANT: John Chicca
                                       ; Sequence 6, Application US/60295722
; GENERAL INFORMATION:
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82.4%;
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82.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 82.4
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 18
SOFTWARE: Patent.pm
SEQ ID NO 6
LENGTH: 47
                                                                              APPLICANT: Barbara Bour
                                                                                                                                                                                                                                                                                                                                                                                                  , ORGANISM: Homo sapiens
US-60-295-722-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TYPE: DNA
; ORGANISM: Homo sapiens
US-60-295-600-6
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Best Local Similarity
Matches 14; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
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Pred. No. 9.4e+03;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 33;
                                                                                                                                                        MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage COMPUTER: HP Vectra 486/33 OPERATING SYSTEM: MSDOS version 6.2 SOFTWARE: ASCII Text
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Havukkala, 11kka J
TITLE OF INVENTION: Polynucleotides isolated from plants
TITLE OF INVENTION: and methods for their use.
FILE REFERENCE: 1054P3
CURRENT APPLICATION NUMBER: US/60/253,378
CURRENT FILING DATE: 2000-11-27
NUMBER OF SEQ ID NOS: 40367
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 31544
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 12.2; DB 6;
Pred. No. 9.1e+03;
0; Mismatches 3;
                  ADDRESSEE: Human Genome Sciences, Inc. STREET: 9410 Key West Avenue CITY: Rockville STATE: Maryland
                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/765,272
FILING DATE: 22-Jan-2001
CLASSIFICATION: CURROWN:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/961,083
FILING DATE: CURROWN:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                NAME: Brookes, A. Anders
REGISTRATION NUMBER: 36, 373
REFERENCE/DOCKET NUMBER: PB340P2
TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8504
TELEFAX: (301) 309-8512
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 330:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 31544, Application US/60253378 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 33 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 330: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    61.0%;
82.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 61.0%;
Best Local Similarity 82.4%;
Matches 14; Conservative
                                                                                                                                      COMPUTER READABLE FORM:
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; ORGANISM: Pinus radiata
US-60-253-378-31544
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 14; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-60-253-378-31544
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TYPE: nucleic acid
STRANDEDNESS: double
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a lake Denison

APPLICANT: Frances Yen
APPLICANT: Bernard Bihain
APPLICANT: Bernard Bihain
APPLICANT: Lydie Bougueleret
TITLE OF INVENTION: GSSP4 Polynucleotides and Polypeptides and Uses Thereof
CURRENT APPLICATION NUMBER: US/60/295,722
CURRENT APPLICATION NUMBER: 2001-06-04
NUMBER OF SEQ ID NOS: 18
SOFTWARE: Patent.pm
SOFTWARE: Patent.pm
SOFTWARE: Patent.pm
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APPLICANT: Mark J. Graham
TITLE OF INVENTION: ANTISENSE MODULATION OF APOLIPOPROTEIN(A) EXPRESSION
FILE REPERRNE: ISPH-0585.
CURRENT APPLICATION NUMBER: US/09/923,515
CURRENT FILING DATE: 2001-08-07
NUMBER OF SEQ ID NOS: 41
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 39
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Choi et. al.
TITLE OF INVENTION: Streptococcus pneumoniae Antigens and Vaccines
NUMBER OF SEQUENCES: 452
CORRESPONDENCE ADDRESS:
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MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage
COMPUTER: HP Vectra 486/33
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STREET: 9410 Key West Avenue
CITY: Rockville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: Antisense Oligonucleotide US-09-923-515-39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            60.0%; Score 12; DB 100.0%; Pred. No. 1.3
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82.4%;
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Best Local Similarity 82.4
Matches 14; Conservative
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Best Local Similarity 100.
Matches 12; Conservative
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COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                ; TYPE: DNA
; ORGANISM: Homo sapiens
US-60-295-722-6
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Sequence 272, Application US/09765272
GENERAL INFORMATION:
APPLICANT: Choi et. al.
TITLE OF INVENTION: Streptococcus pneumoniae Antigens and Vaccines
NUMBER OF SEQUENCES: 452
CORRESPONDENCE ADDRESS:
ADDRESSE: Human Genome Sciences, Inc.
STREET: 9410 Key West Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
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APPLICATION NUMBER: US/09/765,272 FILING DATE: 22-3an-2001 CLASSIFICATION: <UNknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.1e+04;
                                                                                         CLASSIFICATION: CLASSIFICATION DATA:
    APPLICATION DATA:
    APPLICATION NUMBER: 08/961,083
    FTLING DATE: 0CT-30-1997
    ATORNEY/AGENT INFORMATION:
    NAME: Michelle S. Marks
    REGISTRATION NUMBER: 41,971
    REGISTRATION NUMBER: PB340P3
    TELECOMMUNICATION INFORMATION:
    TELEPHONE: (301) 309-8504
    TELEFAX: (301) 309-8512
    INFORMATION FOR SEQ ID NO: 272:
    SEQUENCE CHARACTERISTICS:
                                               APPLICATION NUMBER: US/09/536,784 FILING DATE: 30-Oct-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: PB340P2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 272:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 12;
Pred. No.
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APPLICATION NUMBER: 08/961,083
FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Brookes, A. Anders
REGISTRATION NUMBER: 36,373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FELEFAX: (301) 309-8512
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 31 base pairs
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LENGTH: 31 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: double
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
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COMPUTER READABLE FORM:
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Best Local Similarity 75.0°
Matches 15; Conservative
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Sequence 33, Application US/09591306

GENERAL INFORMATION:
APPLICAT: Massachesetts general Hospital
TITLE OF INVENTION: Novel transcription factor and uses therefor
FILE REFERENCE: 17633/1170
CURRENT APPLICATION NUMBER: US/09/591,306
CURRENT FILING DATE: 2000-06-09
PRIOR PILING DATE: 1996-11-19
NUMBER OF SEQ ID NOS: 41
SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     59.0%; Score 11.8; DB 7; Length 37; 86.7%; Pred. No. 1.5e+04; ive 0; Mismatches 2; Indels
                                                                                                                            ZIP: 20850
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS version 6.2
SOFTWARE: ASCII Text
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          60.0%; Score 12; DB 6; Length 34; 75.0%; Pred. No. 1.1e+04;
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                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/765,272
FILING DATE: 22-Jan-2001
CLASSIFICATION: <UNKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Brookes, A. Anders
REGISTRATION NUMBER: 36,373
REFERENCE/DOCKET NUMBER: PB340P2
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                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/961,083
FILING DATE: <UNKNOWN>
                                           STREET: 9410 Key West Avenue CITY: Rockville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COCATION: (1)..(37)
COTHER INFORMATION: Synthetic primer US-09-591-306-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (301) 309-8504
(301) 309-8512
                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 34 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 316:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: double
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Best Local Similarity 75.0°
Matches 15; Conservative
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Best Local Similarity 86.7
Matches 13; Conservative
                                                                                      STATE: Maryland COUNTRY: USA
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US-09-591-306-33
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LENGTH: 37
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US-09-765-272-316/c
; Sequence 316, Application US/09765272
; GENERAL INFORMATION:
; APPLICANT: Choi et. al.
; TITLE OF INVENTION: Streptococcus pneumoniae Antigens and Vaccines
; NUMBER OF SEQUENCES: 452
                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Choi et. al.
TITLE OF INVENTION: Streptococcus pneumoniae Antigens and Vaccines
NUMBER OF SEQUENCES: 452
CORRESPONDENCE ADDRESS:
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                                                                                                          Length 31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage COMPUTER: HP Vectra 486/33 OPERATING SYSTEM: MSDOS version 6.2 SOFTWARE: ASCII Text
                                                                                                        Query Match 60.0%; Score 12; DB 6; Le Best Local Similarity 75.0%; Pred. No. 1.1e+04; Matches 15; Conservative 0; Mismatches 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Human Genome Sciences, Inc.
STREET: 9410 Key West Avenue
CITY: Rockville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/536,784
FILING DATE: 30-Oct-1997
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION DATE:
FILING DATE: OCT-30-1997
                  ; SEQUENCE DESCRIPTION: SEQ ID NO: 272: US-09-765-272-272
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NAME: Michelle S. Marks
REGISTRATION NUMBER: 41,971
                                                                                                                                                                                                                                                                                                                                               Sequence 316, Application US/09536784 GENERAL INFORMATION:
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TELEPHONE: (301) 309-8504
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TELEFAX: (301) 309-8512
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 20850
COMPUTER READABLE FORM:
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: Maryland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
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Best Local Similarity
'-h-es 15; Conserve
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Search completed: October 2, 2001, 16:55:06 Job time: 17669 sec

HIV-1 gp41 C-helic Probe for human PG Rho G antisense ph

Polymorphic fragme PCR primer ZC17157 Primer for 564/5/6 Mouse vascular end Sequence of forwar

AAA04346

AAX26801

Mycoplasma second

Run on:

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New antisense compounds for inhibiting focal adhesion kinase expression, especially useful for inhibiting retinal neovascularization, or for diagnosing and treating e.g. colon cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; focal adhesion kinase; FAK; signal transduction; cancer;
embryonic development disorder; angiogenic disorder; wound healing;
antisense; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                Human focal adhesion kinase antisense sequence #21
                                                                                                                                                                                                         AAQ50767
AAQ83139
AAQ25250
                                                                                                                                                                                                                                                                                   AAZ47143
AAX38989
                                                                                                                                                                    AAT31052
AAT79727
                                                                                                                                                                                       AAX00053
AAA65581
                                                                         AAQ40032
                                                                                                                                                  AAA65140
         AAZ01125
                                               AAD01093
                                                                 AAX88971
                                                                                                     AAF52006
                                                                                                                                                             AAT03859
                                                                                                                                                                                                                                     AAD02858
                                                                                                                                                                                                                                               AAQ47793
                                                                                                                                                                                                                                                       AAC92387
                                                                                                                                                                                                                                                                AA257915
                                                                                                                                                                                                                                                                                                                                                                                         BP.
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AAC65555 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
 (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Monia BP, Gaarde WA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-006141/01.
19-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-AUG-1999;
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                                                                                            111.8
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RESULT
                                                                                                                                                                                                                                                                                                                                                                                         PCR primer used to Oligonucleotide ca Primer murf4. Syn PCR primer used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human NKCC2 gene e
PCR primer used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        murF2. Syn
primer used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human focal adhesi
Human focal adhesi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oligonucleotide sf
                                                                        (without alignments)
10.740 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Primer murF2
                                                                Search time 876.95 Seconds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Description
                                                                                                                                                                                                                                                                                                 / SIDSB/gcgdata/geneseq/geneseqn/NA1982 DAT: *
/ SIDSB/gcgdata/geneseq/geneseqn/NA1981 DAT: *
/ SIDSB/gcgdata/geneseq/geneseqn/NA1991 DAT: *
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/ SIDSB/gcgdata/geneseq/geneseqn/NA1992 DAT: *
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/ SIDSB/gcgdata/geneseq/geneseqn/NA1992 DAT: *
/ SIDSB/gcgdata/geneseq/geneseqn/NA1994 DAT: *
/ SIDSB/gcgdata/geneseq/geneseqn/NA1995 DAT: *
/ SIDSB/gcgdata/geneseq/geneseqn/NA1999 DAT: *
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR
                                                                                                                                                                                                                                                                                  /SIDS8/gcgdata/geneseq/geneseqn/NA1980.DAT:*/SIDS8/gcgdata/geneseq/geneseqn/NA1981.DAT:*
         GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                    Fotal number of hits satisfying chosen parameters:
                                                                                                                                                                   730101 segs, 313950809 residues
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Maximum Match 100%
Listing first 45 summaries
                                                               2, 2001, 16:18:46
                                            - nucleic search, using sw model
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AAC6535
AAC83406
AAV45725
AAX36171
AAT63364
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AAX86099
AAQ04938
AAT63369
AAX86104
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Gapop 10.0 , Gapext 1.0
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Match Length
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Perfect score:
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Maximum DB s
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Database

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Result Š.

Human fibroblast g Beta-lactoglobulin Human FGFR1-P1 PCR Murine proteinase ERM HIV target seq HIVOV12 No. 1960 NANBH PCR primer P Human GPVI externa

CD100 PCR primer S Escherichia coli t Anabaena variabili

Bilirubin oxidase CD100 PCR primer (

Human genomic DNA Human A-Raf substr Human CYP3A relate

AAV92548

ALIGNMENTS

Liposome membrane

Human integrin alp Measles virus L pr Primer specific fo Primer #3 targeted Primer BLGAMP4 for

Forward primer 34F IGF-I oligonucleot IGF-I oligonucleot

Granule bound star

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The present invention relates to a translational reporter vector selected from p2luc and p2luci. The vectors have a polylinker interposed between a remilla luciferase gene and a firefly luciferase gene, which are out-of-frame with respect to each other but are co-expressed upon recoding. The translational reporter vector is useful for assaying translational recoding in vitro or in vivo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Translational reporter vector for quantification of translation recoding in vivo and in vitro, comprises renilla luciferase gene, polylinker for insertion of selected DNA and an out-of-frame firefly luciferase gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Na-K-2Cl cotransporter; NKCC2; human; Bartter's syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ion transport; hypokalaemic alkalosis; hypercalciuria; nephrocalcinosis; diagnosis; therapy; SSCP; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human NKCC2 gene exon 24 forward primer hNKCC2ex24
                                                                                                                                                                                                                                                                                  reporter vector; renilla luciferase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 13.4; DB 22;
Pred. No. 1.7e+02;
); Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 31 BP; 4 A; 10 C; 9 G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gesteland RF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 4; Column 33; 21pp; English.
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0
                                                                                                                                    BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            89.3%;
93.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  99US-0282996.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (UTAH ) UNIV UTAH RES FOUND.
                                                                                                                                                                                                                                                                                                        recoding; ss.
                                                                                                                                AAC83406 standard; DNA; 31
                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Atkins JF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14; Conservative
 1 gegggeteacagtgg 15
                   3 gcgggctcacagtgg 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 gegggetcacagtgg 15
                                                                                                                                                                                                                                               Oligonucleotide sfAZ2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                    Translational translational
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Grentzmann G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         31-MAR-1999;
                                                                                                                                                                                                          26-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                              US6143502-A.
                                                                                                                                                                                                                                                                                                                                                                                                               07-NOV-2000
                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                       AAC83406;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAV45725;
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                                                                                         RESULT
AAC83406/
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                           The present invention describes a number of phosphorothicate antisense sequences to the human focal adhesion kinase (FAK) protein. This protein is involved in integrin-mediated signal transduction, and is implicated in cancer, particularly colon, breast and oral tumours, embryonic development disorders, angiogenic disorders and wound healing. The antisense sequences, including the one shown here, can be used in the treatment of all of these.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes a number of phosphorothioate antisense sequences to the human focal adhesion kinase (FAK) protein. This protein is involved in integrin-mediated signal transduction, and is implicated in cancer, particularly colon, breast and oral tumours, embryonic development disorders, angiogenic disorders and wound healing. The antisense sequences, including the one shown here, can be used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense compounds for inhibiting focal adhesion kinase expression, especially useful for inhibiting retinal neovascularization, or for diagnosing and treating e.g. colon cancer
                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; focal adhesion kinase; FAK; signal transduction; cancer;
embryonic development disorder; anglogenic disorder; wound healing;
antisense; phosphorothioate; ss.
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                                                                                                                                                                                                                                               22; Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 15; DB 22; Length 20; 100.0%; Pred. No. 22;
                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human focal adhesion kinase antisense sequence #1.
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0
                                                                                                                                                                                   Sequence 15 BP; 2 A; 4 C; 7 G; 2 T; 0 other;
                                                                                                                                                                                                                                               DB
22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 2 A; 7 C; 8 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                               Score 15;
Pred. No.
Claim 3; Column 25; 30pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 3; Column 23; 30pp; English.
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                                                                                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                             100.0%;
100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antisense sequences, inclu
treatment of all of these.
                                                                                                                                                                                                                                          Query Match 100.
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                      gegggeteacagtgg 15
                                                                                                                                                                                                                                                                                                                                                         1 gegggeteacagtgg 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaarde WA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-006141/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-AUG-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US6133031-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAC65535;
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Gaps

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Indels

Length 31;

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Primers murFl (AAT63363) and murF2 (AAT63364) were used to amplify the Escherichia coli murF gene (see also AAT63360) and were designed so as to allow in-frame fusion of the murF coding sequence to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                               The specification describes an intercellular adhesion molecule (ICAM)-6 polypeptide. The ICAM-6 polypeptides and polymelocities can be used for drug screening and developing products for therapy involving intercellular adhesion, e.g. in inflammatory processes. The products can also be used for detection, diagnosis and the production of transgenic animals. PCR primers AAX36170-73 were used in the course of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Culture system for stable high yield production of recombinant plasmids - using bacteria with modified chromosome that can survive only when plasmid is present and optionally foreign DNA expressible only in eukaryote(s)
                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                        New isolated intercellular adhesion molecule-6 used for, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                       Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Recombinant plasmid; DNA immunisation; genetic vaccine; gene therapy; murF gene; Escherichia coli; primer; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                     Score 11.8; DB 20;
Pred. No. 1.2e+03;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                Sequence 24 BP; 6 A; 3 C; 10 G; 5 T; 0 other;
                                                   Vazeau R;
                                                                                                                                                                               Example 15; Page 53; 102pp; English.
                                                                                                                                           diagnosis of inflammatory processes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 20; 46pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP.
                                                                                                                                                                                                                                                                                                                                                                                                                     78.78;
86.78;
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95US-0548059.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                   Staunton DE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 gcgggctcacagtgg 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gctggatcacagtgg 24
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                                                                                     WPI; 1999-288308/24
                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
               (ICOS-) ICOS CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Primer murF2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9714805-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-NOV-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-OCT-1995;
                                                   Loughney K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-APR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                         13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Morsey MA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAT63364;
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                            Thiazide sensitive cotransporter, ATP sensitive K channel and NaK2C1 cotransporter genes - useful for developing products for the diagnosis and treatment of ion transport disorders, e.g. Gitelman's Syndrome or Bartter's Syndrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Primers hNKCC2ex24 forward and reverse (see AAV45725 and AAV45726, respectively) are designed to amplify exon 24 of the human NKCC2 gene (see AAV40562) that codes for Ne K-21 cotransporter NKCC2 protein (see AAV43683). Both primers are located within introns of the gene. 27 Sets of specific primers (see AAV45677-V45730) were used for SSCP analysis of NKCC2. Amplified products were analysed for molecular variants by electrophoresis, and identified variants was demonstrated. Identification of the molecular basis of Bartter's syndrome with NKCC2 Bartter's syndrome allows for the genetic diagnossis of this disorder. The invention provides products and methods useful for diagnosis and treatment of Bartter's syndrome and other ion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Intercellular adhesion molecule 6; ICAM-6; drug screening; therapy; intercellular adhesion; inflammatory process; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR primer used to amplify a fragment of ICAM-6 nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 11.8; DB 19;
Pred. No. 1.2e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                         Example 2; Page 65; 105pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      78.7%;
86.7%;
                                                                                                                                           97WO-US23553
                                                                                                                                                                               96us-0778052.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 86.7
Matches 13; Conservative
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                                                                                                                                                                                                                                                                                        WPI; 1998-388029/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     transport disorders.
                                                                                                                                                                                                                   (UYYA ) UNIV YALE.
                                  Homo sapiens
                                                                  WO9829431-A1
                                                                                                                                                                             31-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-OCT-1997;
                                                                                                                                           19-DEC-1997;
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                                                                                                      09-JUL-1998
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                                                                                                                                                                                                                                                     Lifton RP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                 Synthetic.
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Query Match

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AAX36171 RESULT

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Mismatches

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Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of the invention
                                                                                                                                                                                                                PCR primer; ss
                                                                                                                                                                                                                                                                                                      16-0CT-1996;
                                                                                                                                                                                                                                                          US5922583-A
                                                                                                                                                                                                                                                                                 13-JUL-1999
                                                                                                                                                                                                                                                                                                                            16-OCT-1996;
                                                                                                                                                                                                                                                                                                                                     17-OCT-1995;
30-NOV-1995;
  13;
                                                                                                                                                                                                                                     Synthetic.
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                                                                                                                          AAX86099;
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Matches
  Matches
                                                                                      AAX86099
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR primers AAV42720-21 were used to amplify human furin cDNA (see AAV42731). The amplified product is used in the retroviral vector of the invention to encode a cell bound prodrug convertese for ablation therapy. The specification describes a gene delivery vehicle which directs expression of a non-immunogenic selectable marker or molecule which is capable of activating a previously inactive compound. Vectors expressing the markers and a heterologous sequence are useful in gene therapy. The vectors can be used to deliver a molecule into a target area where it may cause the activation of a previously inactive substance.
                                                                                                                Gaps
murE promoter. The murE promoter was subsequently amplified (see also AAT63185-66). An E. coli strain was constructed with an irreversible alteration (deletion) in its murF gene. This can be used for large-scale prodn. of recombinant plasmids.
                                                                                                                                                                                                                                                                                                   Retroviral vector; gene delivery vehicle; expression; PCR primer; non-immunogenic selectable marker; gene therapy; activation; human; furin; ablation therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Non:immunogenic pro:drug activating enzyme(s) and selectable marker(s) - are used in gene therapy for the treatment of a wide variety of disorder(s)
                                                                                                              .;
0
                                                                                         Length 31;
                                                                                                              Indels
                                                                                        Score 11.8; DB 18;
Pred. No. 1.2e+03;

    PCR primer used to amplify human furin cDNA.

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                                                      Sequence 31 BP; 6 A; 9 C; 13 G; 3 T; 0 other;
                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 8; Page 53; 121pp; English.
                                                                                                                                                                                                               BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Moore MD;
                                                                                        78.78;
86.78;
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97US-0035473.
97US-0038339.
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                                                                                                                                                                                                              AAV42721 standard; DNA; 31
                                                                                                                                                                                                                                                         (first entry)
                                                                                        Ouery Match 78.7
Best Local Similarity 86.7
Matches 13; Conservative
                                                                                                                                   1 gcgggctcacagtgg 15
                                                                                                                                                12 gcaggctgacagtgg 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Jolly DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1998-399153/34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CHAD/) CHADA S.
(JOLL/) JOLLY D J.
(MOOR/) MOORE M D.
                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                WO9830709-A2
                                                                                                                                                                                                                                                                                                                                                                                                                             14-JAN-1998;
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27-FEB-1997;
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                                                                                                                                                                                                                                                        14-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                       16-JUL-1998
                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                    AAV42721;
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                                                                                                                                                                                         RESULT
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Length 31;

Score 11.8; DB 19; Pred. No. 1.2e+03;

78.78; 86.78;

Query Match Best Local Similarity

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                                                                                                                                                                                                                                                                                                                                                      Culture system; high level production; recombinant plasmid; murF; plasmid replication; plasmid production; DNA immunisation; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The specification describes a culture system for stable high-level production of recombinant plasmids. The method comprises propagating irreversibly altered bacterial cells under conditions such that cell viability is dependent on the recombinant plasmid. The system is used is for the replication and high-level production of structurally and genetically stable recombinant plasmids carrying DNA, this DNA can then be used in DNA immunisation or gene therapy. PCR primers AAX86098-99 were used to amplify the murF gene, for use in the method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Culture system for stable and high-level production of DNA contained on recombinant plasmids, with improved plasmid stability
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ó
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 31;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 11.8; DB 20 Pred. No. 1.2e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 31 BP; 6 A; 9 C; 13 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                          PCR primer used to amplify the murF gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Column 13; 27pp; English.
                                                                                                                                                                    BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP.
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95US-0548059.
95US-0564973.
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86.78;
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                                                                                                                                                               AAX86099 standard; DNA; 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA; 39
                                                                                                                                                                                                                                                             13-SEP-1999 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 gcgggctcacagtgg 15
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1 gcgggctcacagtgg 15
                                            13 gtgggctcacagagg
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
nes 13; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ04938;
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mutation

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Culture system; high level production; recombinant plasmid; murF; plasmid replication; plasmid production; DNA immunisation; gene therapy;
                                                                                                                                                                                                                             temperature-sensitive murF gene (see also AAT6336) were used to amplify th temperature-sensitive murF gene (see also AAT6336) from Escherichia (coli TKL-68. The amplified nucleotide sequence was cloned into the BamHI site of pUC19, and can be used to produce temperature-sensitive bacterial host cells useful for large-scale prodn. of recombinant placemide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Culture system for stable and high-level production of DNA contained on recombinant plasmids, with improved plasmid stability
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The specification describes a culture system for stable high-level production of recombinant plasmids. The method comprises propagating irreversibly altered bacterial cells under conditions such that cell
                                                                                                            Culture system for stable high yield production of recombinant plasmids . using bacteria with modified chromosome that can survive only when plasmid is present and optionally foreign DNA expressible only in eukaryote(s)
                                                                                                                                                                                                                                                                                                                                                                                                    Length 39;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                 Score 11.8; DB 18;
Pred. No. 1.3e+03;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                Sequence 39 BP; 9 A; 10 C; 15 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer used to amplify the murF gene.
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                                                                                                                                                                                               Example 2; Page 23; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                    ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP.
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95US-0548059.
95US-0564973.
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86.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 gcgggctcacagtgg 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20 gcaggctgacagtgg 34
                                                                                                                                                                                                                                                                                                                  recombinant plasmids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-404463/34.
             (BIOS-) BIOSTAR INC.
                                                                              WPI; 1997-245120/22
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                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-OCT-1996;
17-OCT-1995;
30-NOV-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    13;
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                                               Morsey MA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX86104;
                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 11
                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Factor VIII analogue is sufficiently mutated from the original gene that it is not recognised by blood Abs of the patient. The analogue is composed of human FVIII but carries the B-domain of human FV in place of the FVIII B-domain.

Oligonucleotides can be used to alter the profile of the blood factor without significantly altering its activity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 11.8; DB 11; Length 39;
Pred. No. 1.3e+03;
); Mismatches 2; Indels (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Recombinant plasmid; DNA immunisation; genetic vaccine; gene therapy; murf gene; Escherichia coli; primer; PCR;
            Oligonucleotide carrying mutation for factor VIII gene.
                                                                                                                                                                                                                                                                                                                                                                                                               New hybrid DNA encoding hybrid procoagulant proteins prepd. by modifying DNA encoding human factor VIII.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 39 BP; 9 A; 11 C; 11 G; 8 T; 0 other;
                                                                                                            Location/Qualifiers
18..23
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                                           Human factor VIII analogue; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gene therapy; murF gene; Esch
polymerase chain reaction; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   78.78;
86.78;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; ; pp; English.
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                                                                                                                                                                                                                                                                                                               (GENE-) GENETICS INST INC
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Best Local Similarity 86.7
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                              Kaufman RJ, Pittman DD;
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                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1990-193265/25
                                                                                                                                                                                                                                              14-NOV-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9714805-A2
                                                                                                                                                                                                                                                                            14-NOV-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Primer murF4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-0CT-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-NOV-1995;
17-OCT-1995;
                                                                                                                                                                              WO9005530-A
                                                                                                                                                                                                               31-MAY-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-SEP-1997
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                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
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RESULT 10 AAT63369

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Gaps

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The invention relates to raising a neutralising antibody response to a broad spectrum of HIV (human immunodeficiency virus) strains and isolates, comprising the administration of a peptide which corresponds to or minics highly conserved portions of the goal envelope glycoprotein which are important in mediating the process of viral entry into host cells. Such peptides can correspond to or minic the coiled coil solution structure of the N-helical domain (the heptad repeat region), or can correspond or minic the C-helical domain (the ransmembrane-proximal amphipathic alpha-helical segment), or the gp41 core 6-helix bundle, which is formed by the interaction of the N- and C-helical domains of three gp41 proteins. The peptides can be administered either singly or as a combination (particularly a combination of N-helical and C-helical peptides), and can be multimerised. For example, N- and C-helical domain peptides can be
                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Raising neutralizing antibody response to human immunodeficiency virus, comprises administering a polypeptide capable of forming a stable coiled-coil solution structure -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HIV-1; gp41; N-helical domain; heptad repeat region; C-helical domain; gp41 transmembrane-proximal amphipathic alpha-helical segment; core 6-helix bundle; viral entry inhibition; immunogenic; antibody; humoral response, broad spectrum vaccine; anti-HIV; envelope glycoprotein; prophylaxis; therapy; PCR primer; ss.
viability is dependent on the recombinant plasmid. The system is used is for the replication and high-level production of structurally and genetically stable recombinant plasmids carriying DNA, this DNA can then be used in DNA immunisation or gene therapy. PCR primers AAX86102-03 were used to amplify the murF gene, for use in the method
                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                        .;
0
                                                                                                                                                                                                 Length 39;
                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                             Score 11.8; DB 20;
Pred. No. 1.3e+03;
0; Mismatches 2;
                                                                                                                                      Sequence 39 BP; 9 A; 10 C; 15 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HIV-1 gp41 C-helical domain 5' PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human immunodeficiency virus type 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 5; Page 54; 97pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                AAA72106 standard; DNA; 44 BP.
                                                                                                                                                                                               78.7%;
86.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-JAN-1999; 99US-0115404
07-JAN-2000; 2000US-0480336.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                .0-JAN-2000; 2000WO-US00456
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                             Query Match 78.7
Best Local Similarity 86.7
Matches 13; Conservative
                                                                                                                                                                                                                                                                           1 gcgggctcacagtgg 15
                                                                                                                                                                                                                                                                                                              20 gcaggctgacagtgg 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-465959/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Weiss CD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (WILD/) WILD C T. (WEIS/) WEISS C D.
                                                                                                    of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200040616-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24 - NOV - 2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA72106;
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                                                                                  and isolates. The peptide compositions may be administered as a prophylactic or therapeutic vaccine to generate antibodies which reduce or inhibit the ability of HIV to infect uninfected cells. A composition comprising polyclonal or monoclonal antibodies can be administered to reduce HIV infection of uninfected cells. Antibodies can be administered to entry-relevant gpd1 structures may also be used therapeutically and as tools to further elucidate the mechanism of HIV cell entry. Sequences AAA72104-A72107 represent PCR primers used in an examplification of the invention to construct DNA (AAA72102) encoding a protein construct (AAB44603) comprising the N- and C-helical domains of HIV-1 gp41 separated by a short peptide linker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PG1 gene; biallelic marker; PCR primer; PG1-related biallelic marker; cancer; prostate cancer; diagnosis; therapy; prostate specific antigen;
alternately linked together to form a peptide which mimics the core 6-helix bundle. Administration of the peptide(s) generates a humoral response, with the production of antibodies against gp41 structures involved in viral entry. As these portions of gp41 are well conserved, such antibodies may be effective against a broad range of HIV strains
                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Use of a prostate cancer associated gene and biallelic markers
                                                                                                                                                                                                                                                                                                                                                                   Length 44;
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                                                                                                                                                                                                                                                                                                                                                                 Score 11.8; DB 21;
Pred. No. 1.3e+03;
                                                                                                                                                                                                                                                                                                            Sequence 44 BP; 13 A; 10 C; 15 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Probe for human PG1 biallelic marker 4-50-293.
                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Blumenfeld M, Bougueleret L, Chumakov I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4; Page 328; 385pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                   0;
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                                                                                                                                                                                                                                                                                                                                                                 78.78;
86.78;
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97US-0996306.
                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 86./v,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ01125 standard; DNA; 47
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                                                                                                                                                                                                                                                                                                                                                                                                                                           15
                                                                                                                                                                                                                                                                                                                                                                                                                                           1 gegggeteacagtgg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           9 gccggctcagagtgg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-405178/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   derived from it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PSA; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAZ01125;
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Gaps

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Length 18; Indels

DB 22; 2e+03;

Score 11.4; DE Pred. No. 2e+03 0; Mismatches

76.0%; 92.3%;

Conservative 14

Similarity

Best Local Sim Matches 12;

Query Match

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pp δλ

2 cgggctcacagtg 17 cegececacaere

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention relates to an antisense compound targeted to a nucleic acid molecule encoding a member of the human Rho family of small GTP binding proteins, where the antisense compound inhibits the expression of the member of the human Rho family. The invention includes antisense oligonucleotides AAF94680 - AAF94637 which target a RhoA nucleotide sequence, AAF94686 - AAF94684 which target a RhoC nucleotide sequence, AAF94769 - AAF94769 which target a Rac 1 nucleotide sequence and AAF94765 - AAF94809 which target a Rac 1 nucleotide sequence and AAF94765 - AAF94809 which target a Cdc42 nucleotide sequence. The antisense compound is useful for treating hyperproliferative conditions, especially cancer, abnormal wound healing or clotting conditions and ischaemia/reperfusion or recoxygenation injury. The compound may also be used to diagnose the above conditions.
             to its inability to discriminate between malignant and non-malignant affections of the organ. A need exists for both a reliable diagnostic procedure which would enable early-stage diagnosis, and for preventative and curative treatments of the disease. The PG1 gene can be used for detection of prostate cancer, and the risk of developing it in the future, and can also be used to determine therapies for the disease.
antigen (PSA) dosage. However, the effectiveness of this is limited due
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rho; GTP binding protein; phosphorothioate antisense oligonucleotide; RhoA; RhoB; RhoC; Rac 1; cdc42; hyperproliferative condition; cancer; wound healing; clotting; ischaemia; reperfusion; reoxygenation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      An antisense compound targeted to a nucleic acid molecule encoding a member of the human Rho family of small GTP binding proteins useful for
                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                Length 47;
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                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                        1.3e+03;
2;
                                                                                                                                                                                                    DB 20;
                                                                                                                                           Sequence 47 BP; 14 A; 9 C; 14 G; 10 T; 0 other;
                                                                                                                                                                                                                   Pred. No. 1.3e; Mismatches
                                                                                                                                                                                                  Score 11.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             e.g. cancer and ischaemia -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 18; Page 81; 156pp; English.
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86.78;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-MAY-2001 (first entry)
                                                                                                                                                                                        Ouery Match
Best Local Similarity 86.7
Matches 13, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cowsert LM;
                                                                                                                                                                                                                                                                                           1 gcgggctcacagtgg 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-191677/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200115739-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          31-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-MAR-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Roberts ML,
                                                                                                                                                                                                                                                                                                                                                                                                                                     AAF94736;
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Polymerase chain reaction; Mycoplasma; M.fermentans; M.hyorhinis; M.arginini; M.pirum; M.hominis; M.salivarium; M.orale; detection; ss.

Mycoplasma second stage PCR primer M34

(first entry)

10-JUL-1997

AAT45334;

AAT45334/c ID AAT45334 standard; DNA; 22 BP.

(AMTY-) AMERICAN TYPE CULTURE COLLECTION.

Roblin RO;

96WO-US07116. 95US-0445289

13-MAY-1996; 19-MAY-1995;

21-NOV-1996

WO9636735-A1.

Synthetic.

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The present sequence is the M34 primer involved in the second stage of pCR in the novel method for amplifying a target segment of nucleic acid in a sample nucleic acid mixture. It is part of the second stage primer mixture that is made up of 3 different primers: F2, R2, and M34 in relative molar ratios of 0.25: 0.125: 1. Using both first and second stage primers in the nested PCR method provides an enhanced system for specific, sensitive and rapid detection of the presence and identification of Mycoplasma species and Mycoplasma-like organism (MLO) contaminants contained in a sample of nucleic acid. The method allows detection of less than 200 colony forming units (cfu)/ml of Mycoplasma species M. Fermentans, M. hyorhinis, M. arginni, M. pirum, M. hominis and M. salivarium and 350 cfu/ml for M. orale. This compares to detection (levels of low of unike prior art Mycoplasma detection systems. Also, unlike prior art Mycoplasma detection systems. The new method and primers can detect the MLO contaminant Acholeplasma laidlawil at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ó
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Two-stage PCR system using sets of mixed and single sequence primers – useful for sensitive detection of mycobacteria and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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llarity 92.3%; Pred. No. 2e+03;
Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 22 BP; 1 A; 8 C; 4 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 18; Page 30; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           levels as low as 20 cfu/ml
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nes 12; Conserv
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Matches
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Sequence 18 BP; 1 A; 6 C; 6 G; 5 T; 0 other;

Search completed: October 2, 2001, 16:18:47 Job time: 15491 sec

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GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
APPLICANT: Garde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310B
CURRENT PILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 42
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence
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   Similarity
   00000000000000
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US-09-377-310-40
SEQ ID NO 40
LENGTH: 15
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
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13, Appl
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Patent No. 5256642
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                                                                                                                      (without alignments)
9.071 Million cell updates/sec
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                                                                                                        Search time 417.38 Seconds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Description
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence Seq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
                              Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-377-310-20
US-08-244 492A-7
US-08-244 492A-7
US-08-244 492A-7
US-08-18-604-5
US-08-18-604-5
US-08-18-604-5
US-08-18-916-14
US-08-18-916-14
US-08-916-18-10
US-08-916-1042
US-08-916-1043
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US-08-916-1043
US-08-916-1043
US-08-916-1043-316
                                                                                                                                                                                                                                                                                                       Total number of hits satisfying chosen parameters:
           GenCore version 4.5
Copyright (c) 1993 - 2000 Comm
                                                                                                                                                                                                                                                                            324599 seqs, 94655562 residues
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                                                                                                      2001, 16:03:53
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Maximum Match 100%
Listing first 45 summaries

    nucleic search, using sw model

                                                                                                                                                                                                  catgatgcttaaaagcttac 20
                                                                                                                                                                                                                               IDENTITY_NUC
Gapop 10.0 , Gapext 1.0
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                                                                                                                                                             US-09-757-100B-20
20
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Match Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Minimum DB seq length: 0 Maximum DB seq length: 50
                                                                                                        October
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                                                                                                                                                                  Title:
Perfect score:
                                                                         OM nucleic
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Sequence 20, Application US/09377310B

Patent No. 6133031

SEREMAL INFORMATION:

APPLICANT: Monia, Brett F.

APPLICANT: Monia, Brett F.

TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase

TITLE OF INVENTION: Expression

TITLE OF INVENTION: Expression

TITLE OF INVENTION: Expression

CURRENT FILING DATE: 1999-08-19

CURRENT FILING DATE: 1999-08-19

NUMBER OF SEQ ID NOS: 43

SOFTWARE: PatentIn Ver. 2.0

SEQ ID NO 20

LENGTH: 20
                                          sednence
            Sequence
                             Sequence
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         US-08-327-451E-29
US-08-458-109-29
US-08-448-180-31
US-08-985-112-1497
US-07-932-905B-84
US-08-700-757-84
US-08-512-811A-7
US-08-511-811A-7
US-08-551-8148-27
US-08-551-344-28
US-08-244-44-8
US-08-828-010-8
US-08-828-010-8
US-08-951-344-3
US-08-951-344-3
US-08-98-98-162-9
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                                                                                                                                                                                                                                                                                                      ALIGNMENTS
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5472939-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CTHER INFORMATION: antisense sequence US-09-377-310-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 40, Application US/09377310B
; Patent No. 6133031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;
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Patent No. 5804417
GENERAL INFORMATION:
APPLICANT: Martens, Gerardus J. M.
APPLICANT: Chaudhuri, Bhabatosh
APPLICANT: Stephan, Christine
TITLE OF INVENTION: Production of Proteins Using 7B2 Protein
NUMBER OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
                                                              /function= "primer for PCR"
/product= "synthetic oligonucleotide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 29;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: CIBA-GEIGY Corporation Patent Department
STREET: 520 White Plains Road P.O. Box 2005
CITY: Tarrytown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRENT AFPLICATION NUMBER: US/08/709,915
FILING DATE: 09-SEP-1996
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/244,492
FILING DATE: 02-SEP-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: NE 91/02009
FILING DATE: 29-NOV-1991
ATTORREY AGENT INFORMATION:
REGISTRATION NUMBER: 33200
REFERENCE/DOCKET NUMBER: 4-19344/A/MAR/PCT/DI
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                               Score 13.8; DB 1;
Pred. No. 2.1e+02;
); Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 13.8; DB 1;
Pred. No. 2.1e+02;
0; Mismatches 2;
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MOLECULE TYPE: DNA (genomic)

US-08-709-915-9
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88.2%;
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88.2%;
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ZIP: 10591-9005
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELERAX: 908-2//
INFORMATION FOR SEQ ID NO: 9
SEQUENCE CHARACTERISTICS:
                   3 tgatgcttaaaagctta 19
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Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
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Matches 15; Conserv
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US-08-709-915-9/c
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                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Martens, Gerardus Julianus Maria
APPLICANT: Chaudhuri, Bhabatosh
APPLICANT: Christine Stephan
TITLE OF INVENTON: Production of Proteins Using 7B2 Protein
CORRESPONDENCE: 13
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PATENTINE CONTROLL OF VERSION #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/244,492A
FILING DATE: 02-September-1994
CLASSIFICATION: 435
PRICA PAPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: PCT/EP92/02740
FILING DATE: 27-NO. 5708140ember-1992
PRICA TON NUMBER: Netherlands 9102009
FILING DATE: 29-NO. 5708140ember-1991
ATTONNEY/AGENT INFORMATION:
                                                                                                     DB 3; Length 15;
54;
                                                                                                                                              0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: SPULIL, W. MUTAN
REGISTRATION NUMBER: 32,943
REFERENCE/DOCKET NUMBER: 4-19344/A/WAR/PCT
TELEPHONE: (919) 541-8615
TELEPHONE: (919) 541-8689
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 29 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MAME/KEY: misc_feature
LOCATION: 3..8
OTHER INFORMATION: /function= "HindIII-site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature
LOCATION: 9..11
OTHER INFORMATION: /function= "stop codon"
                                                                                                 75.0%; Score 15; DB 100.0%; Pred. No. 54; Live 0; Mismatches
        ; OTHER INFORMATION: antisense sequence US-09-377-310-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: CIEA-GEIGY Corporation
STREET: P. O. Box 12257
CITY: Research Triangle Park
STATE: No. 5708140th Carolina
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CQMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                         US-08-244-492A-7/c
; Sequence 7, Application US/08244492A
; Patent No. 5708140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: misc_feature
LOCATION: 12..29
OTHER INFORMATION: /fun
OTHER INFORMATION: codi
                                                                                                   Query Match 75.0
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                  linear
                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
FEATURE:
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APPLICANT: Yellon, Dale
APPLICANT: Yellon, Dale
APPLICANT: Yellon
APPLICANT: Huse, William
APPLICANT: Rosok, Mae J.
TITLE OF INVENTION: No. 5792456el Mutant BR96 Antibodies and
TITLE OF INVENTION: Functional Equivalents Reactive With Human Carcinomas
NUMBER OF SEQUENCES: 62
CORRESPONDENCE ADDRESS:
ADDRESSES: Merchant & Gould
STREDT: 11150 Santa Monica Bldv., Suite 400
                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/487,860
FILING DATE: 07-JUN-1995
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Adriano, Salan B.
REGISTRATION NUMBER: 34,470
REGISTRATION NUMBER: 34,470
REFERENCE/DOCKET NUMBER: 30436.16US01
                                                                                                                                                                                                                                                                                                      Score 13.2; DB 1;
Pred. No. 4.1e+02;
0; Mismatches 3;
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Pred. No. 4.1e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 14, Application US/08487860
Patent No. 5792456
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
TELEPHONE: 310-445-1140
TELEPAX: 310-445-9031
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; MOLECULE TYPE: DNA (genomic)
US-08-487-860-14
                                                                                                                                                                                                            ; MOLECULE TYPE: DNA (genomic)
US-08-285-936-14
TELEPHONE: 310-445-9031
TELEFAX: 310-445-9031
INFORMATION FOR SEQ ID NO: 14
SEQUENCE CHARACTERISTICS:
**PMGTH: 33 base pairs
                                                                                                                                                                                                                                                                                                      66.0%;
83.3%;
                                                                                                                                                                                                                                                                                                                                                                                              3 tgatgcttaaaagcttac 20
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Best Local Similarity 83.3
Matches 15; Conservative
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
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Best Local Similarity
Matches 15; Conserva
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ZIP: 90025-3395
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; . Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Thastrup, Ole
APPLICANT: Tullin, Soren
APPLICANT: Pullin, Soren
APPLICANT: Pullin, Soren
APPLICANT: Pullin, Soren
APPLICANT: Pollien, Lars
APPLICANT: Bjorn, Sara
TITLE OF INVENTION: A Method Of Detecting Biologically
TITLE OF INVENTION: A METHOD OF OR OFFICE
CURRENT APPLICATION NUMBER: 0110/95
EARLIER FILING DATE: 1995-01-31
SEARLIER FILING DATE: 1995-01-31
NUMBER OF SEQ ID NOS: 37
SEGTANDE: FastSEQ for Windows Version 3.0
SEQ ID NO 5:
LENGTH: 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 13.4; DB 2;
Pred. No. 3.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Merchant & Gould
STREET: 11150 Santa Monica Bldv., Suite 400
CITY: Los Angeles
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Adriano, Sarah B. REGISTRATION UNBARE: 34,470
REFERENCE/DOCKET NUMBER: 30436.16USO1
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/285,936 FILING DATE: 04-AUG-1994
                                                                                                               Sequence 5, Application US/08818604C Patent No. 5958713
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-285-936-14/c
; Sequence 14, Application US/08285936
; Patent No. 5728821
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ;
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93.3%;
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; TYPE: DNA
; ORGANISM: Aequorea victoria
US-08-818-604-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Yelton, Dale
APPLICANT: Glaser, Scott
APPLICANT: Huse, William
APPLICANT: Rosok, Mae J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 67.0
Best Local Similarity 93.3
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
ZIP: 90025:3395
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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NUMBER OF SEQUENCES:
                                                                        Houston
                                                                                     STATE: Texas
COUNTRY: US,
ZIP: 77210
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 20850
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                                                                                                                                        Flavia M. MODIFIED RECEPTORS THAT CONTINUOUSLY
                                                                                                                                                                                                                                                                                                  COUNTRY: USA

ZIP: 77210-4433

COMPUTER READABLE FORM:
MEDIUW TYPE: FILIPOPPY disk
COMPUTER: IBM PC COMPATIBLE
OPPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/762,308
FILING DATE: 09-DEC-1996
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/224,593
FILING DATE: 05-APR-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: United States of America POSTAL CODE: 78701
TELEPHONE NO: (512) 499-4462
TELEFAX: (512) 499-453
VENTION: MODIFIED RECEPTORS THAT VENTION: CONTINUOUSLY SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         201 West 7th Street
Austin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Kitchell, Barbara S.
REGISTRATION NUMBER: 33,928
REFERNEC/DOCKET NUMBER: UTSD:335--1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 418-3000
                                                                                                  GENERAL INFORMATION:
APPLICANT: Beutler, Bruce A.
APPLICANT: Bazzoni, Flavia M.
TITLE OF INVENTION: MODIFED RECEPTORE:
TITLE OF INVENTION: SIGNAL
CORRESPONDENCE ADDRESS:
ADDRESSEE: Arnold White & Durkee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Sequence 5, Application PC/TUS9504221
; GENERAL INFORMATION:
; APPLICANT:
                                                                  Sequence 5, Application US/08762308
Patent No. 5925548
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
US-08-762-308-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            66.0%;
83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 5
SEQUENCE CHARACTERISTICS:
LENGTH: 33 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Texas
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Matches 15; Conservative
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                                                                                                                                                                                                                                               STREET: P.O. Box 4433
CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE:
COUNTRY:
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STREET:
CITY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 9
PCT-US95-04221-5/c
                                                                                                                                                                                                                                                                     CITY: Hous
STATE: TX
                                                 US-08-762-308-5/c
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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT:
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Sequence 4.18, Application US/08961083
Patent No. 6159469
GENERAL INFORMATION:
TITLE OF INVENTION: Streptococcus pneumoniae Antigens and Vaccines NUMBER OF SEQUENCES: 452
CORRESPONDENCE ADDRESS:
ADDRESSEE: Human Genome Sciences, Inc.
STREET: 9410 Key West Avenue
CITY: Rockville
STATE: Maryland
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 13.2; DB 5;
Pred. No. 4.1e+02;
0; Mismatches 3;
                                                                                                                                                                                                                     OPERATING STSTEM: PC-DOS/MS-DOS/ASCII SOFTWARE: WordPerfect 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: PCT/US95/04221
FILING DATE: CONCURRENTLY HEREWITH
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US SN 08/224,593
FILING DATE: 05 APRIL 1994
CLASSIFICATION:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Arnold, White & Durkee
STREET: P.O. Box 4433
                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: KITCHELL, BARBARA S.
REGISTRATION NUMBER: 33,928
REPERENCE/DOCKET NUMBER: UTT
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (512) 418-3000
TELEPAX: (713) 789-2679
TELEX: 79-0924
INFORMATION FOR SEQ ID NO: 5:
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83.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 66.0
Best Local Similarity 83.3
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 33 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      , MOLECULE TYPE: DNA PCT-US95-04221-5
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GENERAL INFORMATION:

APPLICANT: SASMI, Keiko

APPLICANT: ARAJUKI

APPLICANT: ARAJUKI

APPLICANT: MORI, TAKAJUKI

APPLICANT: MAKINO, SATOSHI

TITLE OF INVENTION: CONTAINING SPECIFIC NUCLEOTIDE SEQUENCE AND A METHOD FOR TITLE OF INVENTION: ITS ABSOLUTE IDENTIFICATION

NUMBER OF SEQUENCES: 19

CORRESPONDENCE ADDRESSE: YOUNG & THOMPSON

STREET: 745 SOUTH 23rd Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/905,817
                                                                                                                                                                                      SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
FILLING DATE: 25-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 12.2; DB 1;
Pred. No. 1.2e+03;
0; Mismatches 3;
                                                                                                                                                                                                                                                                     CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/848,400
FILING DATE: 10-MAR-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 3-293625
FILING DATE: 14-00T-1991
ATORNEY AGENT INFORMATION:
NAME: PATCH, Andrew J.
REGISTRATION NUMBER: 32,925
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        KP-7501
                                                                                                  COMPUTER READBLE FORM:
BDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 10, Application US/08905817; Patent No. 5824777
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: Floppy disk
IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: KF
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-521-2297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         61.0%;
82.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 61.0
Best Local Similarity 82.4
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 catgatgcttaaaagct 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: 703-52-2-70573 - 703-685-0573
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 25 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25 CATGATGCTTGAGAGGT 9
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
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                                              Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Arlington
                     Arlington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
TOPOLOGY: lin
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US-08-905-817-10/c
                                                                                   22202
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                                                               COUNTRY:
                                         STATE:
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APPLICANT: MORI, Takayuki
APPLICANT: MAKINO, Satoshi
TITLE OF INVENTION: ATTENDATED MEASLES VIRUS VACCINE,
TITLE OF INVENTION: CONTAINING SPECIFIC NUCLEOTIDE SEQUENCE AND A METHOD FOR
TITLE OF INVENTION: ITS ABSOLUTE IDENTIFICATION
NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ó
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Robert McKay
APPLICANT: Madeline M. Butler
APPLICANT: Madeline M. Butler
APPLICANT: Jacqueline Wyatt
APPLICANT: Lex M. COWSETT
TILLE OF INVENTION: ANTISENSE MODULATION OF PEPCK-CYTOSOLIC EXPRESSION
FILE REFERENCE: RTS-0123
CURRENT APPLICATION NUMBER: US/09/488,671A
CURRENT FILING DATE: 2000-01-21
NUMBER OF SEQ ID NOS: 177
SEQ ID NO 75
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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Pred. No. 9.7e+02;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                   Length 36;
                                                                                                                                                                                                                                                                                                                                                                              Score 12.6; DB 3;
Pred. No. 8.1e+02;
0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ); OTHER INFORMATION: Antisense Oligonucleotide US-09-488-671-75
                                   ATTORNEY/AGENT INFORMATION:
NAME: Brookes, A. Anders
REGISTRATION NUMBER: 36,373
REFERENCE/DOCKET NUMBER: PB340P2
TELECOMMUNICATION INFORMATION:
TELEPAX: (301) 309-8504
TELEFAX: (301) 309-8512
INFORMATION FOR SEQ ID NO: 418:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 75, Application US/09488671A Patent No. 6187545 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 10, Application US/08348891A Patent No. 5654136 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          YOUNG & THOMPSON
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78.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 62.0%;
Best Local Similarity 92.9%;
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2 atgatgcttaaaagcttac 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21 ATTATGTTGATAAGCTTAC 3
                                                                                                                                                                                                                                                                                                                                                                              Query Match 63.0
Best Local Similarity 78.9
Matches 15; Conservative
                                                                                                                                                                                                                                LENGTH: 36 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4 gatgettaaaaget 17
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                               linear
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                                                                                                                                                                                                                                                                                             ; TOPOLOGY:
US-08-961-083-418
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-488-671-75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
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APPLICANT: RUBEN, STEVEN M
APPLICANT: SANCAR, AZIZ
APPLICANT: HSU, SHIAO-WEN D
APPLICANT: HSU, SHIAO-WEN D
APPLICANT: AZANTSEV, ALEKSEY G
TITLE OF INVENTION: HUMAN BLUE-LIGHT PHOTORECEPTOR hCRY2
NUMBER OF SEQUENCES: 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                   The letter "N" stands for the stem
II region of a HH ribozyme.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX P.L.L.C.
STREET: 1100 NEW YORK AVENUE, NW, SUITE 600
CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTY: USA

ZIP: 20005-3934

COMPUTER READABLE FORM:
MEDIOW TYPE: FLOPPY disk
COMPUTER: IBM PC COMPATIBLE
OFFRATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/964,268
FLING APPLICATION DATA:
APPLICATION NUMBER: US 60/030,189
FLING APPLICATION NUMBER: US 60/030,189
FLING DATE: 04 NOV-1996
ATTOREY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 12.2; DB 3;
Pred. No. 1.2e+03;
4; Mismatches 4;
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                                              230/107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 13, Application US/08964268
; Patent No. 6114503
                                       REFERENCE/DOCKET NUMBER: 230, TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 955-0440 TELEFAX: (213) 955-0440 TELEX: 67-3510 INFORMATION FOR SEQ ID NO: 1042:
                         REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              36,688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RECISTRATION NUMBER: 36,688
REFERENCE/DOCKET NUMBER: 14
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                 61.0%;
55.6%;
  Warburg, Richard J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3 tgatgcttaaaagcttac 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10 UGAUGANGAAAAGCUUGC 27
                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 55.6
Matches 10; Conservative
                                                                                                                                                                             SEQUENCE CHARACTERISTICS
LENGTH: 27 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 202-371-2540 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: STEFFE, ERIC K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                202-371-2540
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
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COTHER INFORMATION:
US-08-985-162-1042
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US-08-964-268-13
                                                                                                                                                                                                                                                                           TOPOLOGY: linear
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Pred. No. 1.2e+03;
0; Mismatches 3; Indels
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APPLICANT: AFLAT, Saghir
APPLICANT: Fell, Patricia
APPLICANT: McSwiggen, James
TITLE OF INVENTION: GENEMATIC NUCLEIC ACID TREATMENT
TITLE OF INVENTION: OF DISEASES OR CONDITIONS RELATED
TITLE OF INVENTION: TO LEVELS OF EPIDERMAL GROWTH
TITLE OF INVENTION: TO LEVELS OF EPIDERMAL GROWTH
TITLE OF INVENTION: FACTOR RECEPTORS
NUMBER OF SEQUENCES: 1877
CORRESPONDENCE ADDRESS:
ADDRESSE: Lyon & Lyon
STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
           CLASSIFICATION: 435
CLASSIFICATION: 435
CLASSIFICATION DATA:
APPLICATION NUMBER: 0S 08/348,891
FILING DATE: 25-NOV-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 0S 07/848,400
FILING DATE: 10-MAR-1992
PRIOR APPLICATION NUMBER: 3P 3-293625
FILING DATE: 14-OCT-1991
ATORNEY/AGENT INFORMATION:
NAME: PATCH, ANGTEW J.
RECISTRATION NUMBER: 32,925
REFERENCE/DOCKET NUMBER: 32,925
REFERENCE/DOCKET NUMBER: 32,925
TELECHOME: 703-521-2297
TELECHOME: 703-652-2537
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ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FASTEDO for Windows 2.0
CURRENT APPLICATION DATA:
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FILING DATE: 04 December 1997
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-985-162-1042; Sequence 1042, Application US/08985162; Patent No. 6057156; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/036,476
FILING DATE: 31 January 1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 10: SEQUENCE: CHARACTERISTICS: LENGTH: 25 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 82.4%;
Matches 14; Conservative
04-AUG-1997
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                                                     Gaps
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()
Ouery Match
61.0%; Score 12.2; DB 3; Length 27;
Best Local Similarity 82.4%; Pred. No. 1.2e+03;
Matches 14; Conservative 0; Mismatches 3; Indels
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Search completed: October 2, 2001, 16:03:53 Job time: 14597 sec

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Mouse A1 T cell re Plasmid pLACII-rec Human HNRCR PCR pr Human flt1 VEGF re Fokt Cleavage doma Human tissue facto Ribozyme vector ol Plasmid insert pGE Ribozyme vector ol Ribozyme flt1 VEGF rec Frimer P4 for anne 3' primer to gener AAV2/4 bottom prim Primer for DNA enc Streptcoccus pneu Prion protein forw Prion protein forw Prion protein forw Prion protein forw Mouse prion protei Synthetic plasmid Human PEPCK-cytoso Corn starch branch cdk4 ribozyme bind Human KDR VEGF rec Human EFR-R mammer Human EFR-R mammer Human BFR-R mammer Human BFR-R memmer Human BFR-R memmer FR-R mammer Human BFR-R memmer FR-R memmer Human BFR-R memmer Human

A STREET

5' 75 kD TNF rece Mouse Al T cell re Plasmid pLAC11-re Human HNRCR PCR p Human fl11 vEGF r FokI cleavage dom Human tissue fact, Ribozyme vector o Plasmid insert pG	RIDOZYME VECTOF O RIDOZYME VECTOF O Human KDR VEGF re Human fill VEGF re Primer P4 for ann 3' primer to gene AAVZ/4 bottom pri Primer for DNA en Streptococcus pnee	Prior protein for Prior protein for Mouse prion protein for Human prion protein Synthetic plasmid Human pepck-cytos Corn starch branc cold fibozyme bin	ribozyme n growth n KDR VEG n fltl VEG n EGF-R h n blue-lin	<pre>tence #18. transduction; cancer; disorder; wound healing;</pre>		focal adhesion kinase oiting retinal
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4.5 Compugen Ltd. ; Search time 876.95 Seconds (without alignments) 14.320 Million cell updates/sec		854978	eqn/Na1980.DaT:* eqn/Na1980.DaT:* eqn/Na1982.DaT:* eqn/Na1983.DaT:* eqn/Na1984.DaT:* eqn/Na1985.DaT:* eqn/Na1986.DaT:* eqn/Na1986.DaT:*	eqn/NA1989. DAT: * eqn/NA1990. DAT: * eqn/NA1991. DAT: * eqn/NA1992. DAT: * eqn/NA1992. DAT: * eqn/NA1995. DAT: * eqn/NA1995. DAT: * eqn/NA1995. DAT: * eqn/NA1996. DAT: * eqn/NA1996. DAT: * eqn/NA1999. DAT: * eqn/NA1999. DAT: * eqn/NA1999. DAT: * eqn/NA1999. DAT: *	cted by chance to have a of the result being printed, score distribution.	Description Human focal adhesi Bacterial 165 rRNA Human focal adhesi Helicobacter pylor Staphylococcus aur Anti-sense strand Primer used to amp cAMP receptor prot Anti-metastatic fu Human serum albumi 3'PCR primer for i
GenCore version 4.5 Copyright (c) 1993 - 2000 Comp nucleic search, using sw model october 2, 2001, 16:18:45; Sea	score: 20-09-757-100B-20 1 catgatgcttaaaagcttac 20 table: IDENTITY_NUC Gapop 10.0 , Gapext 1.0 3: 730101 seqs, 313950809 residues	·	N_Geneseq_0601:* 1: SIDSB/gcgdata/geneseq/genese 2: SIDSB/gcgdata/geneseq/genese 3: SIDSB/gcgdata/geneseq/genese 4: SIDSB/gcgdata/geneseq/genese 5: SIDSB/gcgdata/geneseq/genese 6: SIDSB/gcgdata/geneseq/genese 7: SIDSB/gcgdata/geneseq/genese 8: SIDSB/gcgdata/geneseq/genese 9: SIDSB/gcgdata/geneseq/genese 9: SIDSB/gcgdata/geneseq/genese	10: /SIDSB/gcgdata/geneseq/geneseqn/NA1989.DAT:* 11: /SIDSB/gcgdata/geneseq/geneseqn/NA1990.DAT:* 12: /SIDSB/gcgdata/geneseq/geneseqn/NA1991.DAT:* 13: /SIDSB/gcgdata/geneseq/geneseqn/NA1992.DAT:* 14: /SIDSB/gcgdata/geneseq/geneseqn/NA1992.DAT:* 15: /SIDSB/gcgdata/geneseq/geneseqn/NA1994.DAT:* 16: /SIDSB/gcgdata/geneseq/geneseqn/NA1994.DAT:* 17: /SIDSB/gcgdata/geneseq/geneseqn/NA1995.DAT:* 18: /SIDSB/gcgdata/geneseq/geneseqn/NA1997.DAT:* 19: /SIDSB/gcgdata/geneseq/geneseqn/NA1997.DAT:* 20: /SIDSB/gcgdata/geneseq/geneseqn/NA1999.DAT:* 21: /SIDSB/gcgdata/geneseq/geneseqn/NA1999.DAT:* 22: /SIDSB/gcgdata/geneseq/geneseqn/NA1999.DAT:*	. No. e grea Is der	core Match Length DB ID 20 100.0 20 22 AAC65552 16.4 82.0 26 22 AAC65552 14.2 71.0 29 21 AAZ8455 13.8 69.0 29 14 AAQ42521 13.6 68.0 48 19 AAV43258 13.4 67.0 44 17 AAT39690 13.2 66.0 20 17 AAT12926 13.2 66.0 20 17 AAT12926 13.2 66.0 20 17 AAT12926
OM nucleic Run on:	Title: Perfect sc Sequence: Scoring ta	Total number of Minimum DB seq Maximum DB seq Post-processing	Database :		Pred. score and ii Result	NO. S

Query Match

8 X C C C C C C C X X 8

AAF81601;

RESULT 2 AAF81601/c

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The present invention describes a number of phosphorothioate antisense sequences to the human focal adhesion kinase (FAK) protein. This protein is involved in integrin-mediated signal transduction, and is implicated in cancer, particularly colon, breast and oral tumours, embryonic development disorders, angiogenic disorders and wound healing. The antisense sequences, including the one shown here, can be used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Neutrophil activating protein; NAP; stomach; ulcer; cancer; neutrophil; activation; adhesion; inflammatory response; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New antisense compounds for inhibiting focal adhesion kinase expression, especially useful for inhibiting retinal necowascularization, or for diagnosing and treating e.g. colon cancer
                                                                                                                                                                                                                                                                                      Human; focal adhesion kinase; FAK; signal transduction; cancer; embryonic development disorder; angiogenic disorder; wound healing; antisense; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Helicobacter pylori neutrophil activating protein (NAP) PCR
                                                                                                                                                                                                                                                      Human focal adhesion kinase antisense sequence #38.
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100.0%; Pic
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                                                                                                                                    AAC65572 standard; DNA; 15
  1 catgatgcttaaaagctt 18
                      18 CATGTTGCTTAAAAGCTT 1
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                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
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AAC65572
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                                The present invention describes a number of phosphorothioate antisense sequences to the human focal adhesion kinase (FAK) protein. This protein is involved in integrin-mediated signal transduction, and is implicated in cancer, particularly colon, breast and oral tumours, embryonic development disorders, angiogenic disorders and wound healing. The antisense sequences, including the one shown here, can be used in the treatment of all of these.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention provides several probes and PCR primers which can be used in the detection of bacteria belonging to the genera of Microbispora, Microtetraspora, Nonmuria and Planobispora. The sequences hybridise to the 165 rRNA coding sequence of the organism. The present sequence is a PCR primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel nucleic acid probe hybridizing to nucleic acid encoding portion of 16S rRNA of bacteria of genera Microbispora, Microtetraspora, Nonomuria and Planobispora, useful for detecting these bacteria in
                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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Pred. No. 0.73;
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                                                                                                                                                                                         Sequence 20 BP; 7 A; 4 C; 3 G; 6 T; 0 other;
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Claim 3; Column 23; 30pp; English.
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                                                                                                                                                                                                                                                                      Best Local Similarity 100.
Matches 20; Conservative
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Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2001-281685/29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Microtetraspora.
Planobispora.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200123608-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Genilloud O,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Microbispora
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-JUN-2001
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Gaps

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Query Match

sample

Length 15; Indels

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The present invention describes Staphylococcus thioredoxin reductase (TrxB). The present sequence represent a PCR primer for S. aureus TrxB. TrxB inhibitors can be used as antimicrobials to treat a Staphylococcus, particularly S. aureus, infection. TrxB inhibitors are antimicrobials to which, unlike most of those in the prior art, Staphylococcus has not yet
                                                                                                                                                                                                                                                                                                                                                                       A new thioredoxin reductase from Staphylococcus aureus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Anti-sense strand of the coding sequence of human 7B2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7B2; chaperone function; deaggregation; PCR; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 13.8; DB 20
Pred. No. 7.2e+02;
); Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 27 BP; 7 A; 6 C; 4 G; 10 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Stephan C;
                                                                                                                                                                                                                                                                                        Cohen G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 complement (9..29)
                                                                                                                                                                                                                                                                                                                                                                                                                    Example 3; Page 32; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STICHTING KATHOLIEKE. GEIGY AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             69.0%;
88.2%;
                                                                                                                                                        99WO-US04512
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     92WO-EP02740
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            91NL-0002009
                                                                                                                                                                                                   98US-0076525
                                                                                                                                                                                                                                                                                      Aharonowitz Y, Borovok I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ42521/c
ID AAQ42521 standard; DNA; 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Martens GJM, Chaudhuri B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 69.0
Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 atgatgcttaaaagctt 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20 ATCAAGCTTAAAAGCTT 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        developed a resistance.
                                                                                                                                                                                                                                                                                                                              WPI; 1999-551044/46
                                                                                                                                                                                                                                           (ABBO ) ABBOTT LAB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1993-197066/24
      Synthetic.
Staphylococcus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (UYKA-) UNIV
(CIBA ) CIBA
                                                                   WO9945123-A1.
                                                                                                                                                                                                   02-MAR-1998;
                                                                                                                                                        02-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-NOV-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-NOV-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-SEP-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9311248-A.
                                                                                                             10-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-JUN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ42521;
        qq
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This sequence represents NAP PCR primer #2, used with primer #1

(AAZ34654) to amplify DNA (AAZ34653) encoding the neutrophil activating
protein (NAP) from Helicobacter pylori strain CCUG. The PCR product
was then cloned into the Escherichia coli and Bacillus subtilis
was then cloned into the Escherichia coli and Bacillus subtilis
shuttle vector pSMZ14G for recombinant production. In humans
infected with H.pylori, NAP promotes activation and adhesion
of neutrophils to endothelial cells, an inflammatory response.
C. Since H. Pylori is responsible for stomach inflammation, it is possible
that NAP elicits the inflammatory response, probably at an early stage
of gastric uncer disease, when neutrophils accumulate in the superficial
castif mucosa. The invention relates to a novel process for enriching
the presence of Helicobacter pylori NAP in a mixture of proteins. This
comprises the steps of salting out other proteins (NAP being soluble at
comprises the steps of salting out other proteins (NAP being soluble at
comprises the steps of salting out other proteins (NAP being soluble at
contrastic mucosa.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (preferably with immobilised nickel cations). The NAP purified or enriched in this way is free from amino acid sequences or alterations introduced during cloining (e.g., polyhistidine tags, thioredoxin fusions or GST fusions). NAP produced via this process may be useful for the production of therapeutic agents, such as vaccines. It may also be useful in an immunoassay to ascertain antibody levels in a body sample,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Enriching presence of Helicobacter pylori neutrophil activating protein, useful in diagnostic and therapeutic products and processes
cloning artefact; salting out; ammonium sulphate; metal chelate chromatography; therapy; diagnosis; therapeutic agent; vaccine; immunoassay; immunodiagnosis; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          71.0%; Score 14.2; DB 21; Length 29; llarity 84.2%; Pred. No. 4.6e+02; Conservative 0; Mismatches 3; Indels (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Staphylococcus; trxB; thioredoxin reductase; antimicrobial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 29 BP; 8 A; 8 C; 4 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Staphylococcus aureus trxB PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 8; 26pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and in immunodiagnosis kits.
                                                                                                                                                                                                                                         99WO-IB00695
                                                                                                                                                                                                                                                                                    98GB-0007721
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 catgatgcttaaaagctta 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23 CATTTGGCTTAAAAGCTTA 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ21085 standard; DNA; 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-NOV-1999 (first entry)
                                                                                     Synthetic.
Helicobacter pylori.
                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-052659/04.
                                                                                                                                                                                                                                                                                                                            (CHIR-) CHIRON SPA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR primer; ss.
                                                                                                                                                                                                                                         07-APR-1999;
                                                                                                                                                                                                                                                                                 08-APR-1998;
                                                                                                                                                    WO9953310-A1
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20; Length 27; Indels

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Uziel O,

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                                                                                           The source of the DNA is partly human genomic DNA whilst the immediate experimental source is synthetic. AAQ42515 and AAQ42521 are used in the construction of a truncated 7B2 gene encoding an amino terminal 7B2 signal peptide and the first 170 AAs of the human protein. They are used as primers in a PCR using complete
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Non-covalent complex; antibody; immunoglobulin; immunogen; vaccine; cytotoxic; expression agent; treatment; cancer; expression; transgene;
                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New non-covalent complex of targetting antibody and immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAV43257-58 represent primers used to amplify plasmid pK5DT. The specification describes a non-covalent complex that comprises at least one antibody, or its fragments, able to bind a cell surface molecule and a component that binds (to only 1 type of site) an immunoglobulin (Ig) or fragment, and is associated with an active substance. The complexes are used as immunogens, vaccines or as cytotoxic or expression agents, e.g. for treatment of cancer or
Efficient expression and de-aggregation of protein - by treating co-expressing protein, e.g. hormone, growth or coagulation factor or neuro-peptide with 7B2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         binding component - linked to active agent, e.g. antigen, drug or
nucleic acid, used for treatment of cancer
                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                         Length 29;
                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                      Score 13.8; DB 14;
Pred. No. 7.2e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 48 BP; 14 A; 9 C; 10 G; 15 T; 0 other;
                                                                                                                                                                                                          Sequence 29 BP; 11 A; 6 C; 3 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (COMS ) COMMISSARIAT ENERGIE ATOMIQUE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Primer used to amplify plasmid pK5DT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 6; Page 20; 42pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ä
                                                               Example; Page 36; 41pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                  BP.
                                                                                                                                                                                                                                                      69.0%;
llarity 88.2%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98WO-FR00227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97FR-0001420
                                                                                                                                                                                                                                                                                                                                                                                                                              AAV43258 standard; DNA; 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               or expressing transgenes.
                                                                                                                                                                                                                                                                                                                      tgatgcttaaaagctta 19
                                                                                                                                                                                                                                                                                                                                     Leonetti M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1998-447169/38.
                                                                                                                                                                                                                                                   Query Match'
Best Local Similarity
Matches 15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-0CT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-AUG-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Drevet P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAV43258;
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                                                                                                                                                                                                                                                                                                                                                                                                   RESULT
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Gaps
                                                                                                                                                                                                                                                                                                                                                  intracellular process; fluorescence; probe; screen; identification; luminescent; real time; study; second messenger; enzyme; primer; protein kinase; PCR; polymerase chain reaction; cAMP; receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A claimed fusion construct containing sequences encoding the Aequorea victoria green fluorescent protein (GFP) and a portion of the CAMP receptor protein, i.e. a sequence PCR amplified using the PCR primer pair AAT39689/30, can be used for the detection of a biologically active substance affecting intracellular processes. A cell containing the construct is cultured under conditions permitting GFP expression, and its fluorescence measured. The cell is then incubated with the substance, and its fluorescence massured, where any change indicates that the substance is phologically active. The construct can also be used to generate probes for use in basic research, and screening programmes to
                                                                                                                                                                                                                                                                                                                                     Green fluorescent protein; detection; active substance; affecting;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         identify new biologically active substances. The use of fundinescent probes allows real time studies of 2nd messengers and specific enzymes, e.g. protein kinases, in single living cells, making possible the study of the precise timing and spatial characteristics of these factors. Due to the strong filucrescence of GFP, the luminescence of cells expressing the probes, which are easily introduced into cells, can be easily detected and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New construct encoding modified green fluorescent protein - conie.g. enzyme recognition site, useful for detecting biologically active substances affecting intracellular processes
                                  ö
Length 48;
                                  Indels
Score 13.6; DB 19;
Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 44 BP; 12 A; 8 C; 11 G; 13 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tullin
                                  Mismatches
                                                                                                                                                                                                                                                                                                  CAMP receptor protein DNA, PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Thastrup O,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 3; Page 20; 48pp; English.
                                                                                                                                                                                              BP.
68.0%;
80.0%;
                                                                   1 catgatgcttaaaagcttac 20
                                                                                                    27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   95DK-0000982.
95DK-0000110.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  96WO-DK00052
                                                                                                                                                                                            AAT39690 standard; DNA; 44
                                                                                       || ||| ||| |||| |||||| || caggatccttataagcttcc
                                                                                                                                                                                                                                                               04-APR-1997 (first entry)
                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (NOVO ) NOVO-NORDISK AS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Poulsen LK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1996-371444/37.
              Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9623898-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-JAN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-JAN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                          protein; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-AUG-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bjorn SP,
                                                                                                                                                                                                                               AAT39690;
 Query Match
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                                                                                                                                                                          AAT39690
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Score 13.4; DB 17; Length 44; Pred. No. 1.2e+03;

67.0%; 93.3%;

Query Match Best Local Similarity

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3'PCR primer for isolation and characterisation of P. pastoris FLD1 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Restriction enzyme recognition sites are introduced into the human serum albumin gene at one or more positions sites are pries preparation of fusion genes. The restriction sites are prefuncaduced at the N-terminus, he C-terminus, between the first and second domains and/or between the second and third domains of HSA. The modified HSA gene can then readily be fused to a sequence coding for a physiologically active peptide, e.g. a cancer metastasis inhibitor. The present sequence is that of a PCR primer which was used in the construction of coding sequences ANTI2919 and ANTI2920 which had restriction sites introduced into the gene at positions corresp to the region between domains 1 and 2 or between domains 2
                                                                                                                                                                                                                                                                                                                                                                      Human serum albumin gene modified by introduction of restriction site – useful for production of fusion proteins by inserting active peptide coding sequence into new restriction site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Methylotrophic yeast; formaldehyde dehydrogenase 1; FLD1; marker; formaldehyde resistance; PCR primer; ss.
                    HSA; modify; mutagenesis; genetic engineering; fusion gene; restriction endonuclease; recognition site; amplification; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 13.2; DB 17;
Pred. No. 1.4e+03;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 6 A; 3 C; 2 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 2 and 3; Page 17; 19pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ50168 standard; DNA; 22 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     66.0%;
83.3%;
                                                                                                                                                                                                                 94JP-0209369.
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                                                                                                                                                                                                                                                                                              (ASAG ) ASAHI GLASS CO LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 atgatgcttaaaagctta 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20 ATGAAGATCAAAAGCTTA 3
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Best Local Similarity 83.3
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                WPI; 1996-174566/18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and 3, respectively
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pichia pastoris.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200001829-A2
                                                                                                                                     JP08051982-A
                                                                                                                                                                                                                 11-AUG-1994;
                                                                                                                                                                                                                                                       11-AUG-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-JUL-1998;
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                                                                                                                                                                           27-FEB-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-JAN-2000
                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAZ50168;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT27602-T27615 are PCR primers used for the construction of vectors carrying human serum albumin (HSA) fusion proteins (FPS). The HSA FPS contain a peptide capable of inhibiting metastasis of cancerous cells. The primers are used to introduce restriction sites into the HSA gene for efficient expression and recovery of the FP. Introduction of the anti-metastatic peptide does not destroy the stereo structure of HSA. Previously such fusion proteins could only be produced by a combination of a chemical synthetic method and a binding method. Using recombinant DNA technology for FP prodn. provides a steady supply of the fusion protein on an industrial scale.
      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fusion protein comprising human serum albumin contg. peptide inserted at arbitrary position - useful for, e.g. inhibiting cancer metastasis
                                                                                                                                                                                                                                                                                        Anti-metastatic fusion protein-encoding vector construction primer.
                                                                                                                                                                                                                                                                                                                            Cancer; metastasis; fusion protein; non-disruptional; inhibitor; arbitrary insertion; human serum albumin; HSA; PCR; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
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Pred. No. 1.4e+03;
0; Mismatches 3; Indels (
    Indels
    1;
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    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human serum albumin gene PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 5; Page 21; 24pp; Japanese.
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                                                                                                                                                       AAT27603/c
ID AAT27603 standard; DNA; 20 BP.
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83.3%;
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ID AAT12926 standard; DNA; 20
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                                                                                                                                                                                                                                                   03-SEP-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20 ATGAAGATCAAAAGCTTA 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 66.0°
Best Local Similarity 83.3°
Matches 15; Conservative
    Conservative
                                      5 atgcttaaaagctta 19
|| || || || || || || || || || || 5
5 attcttaaaagctta 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1996-175732/18.
                                                                                                                                                                                                                                                                                                                                                                                                                                               JP08053500-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-AUG-1994;
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14;
                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                               AAT27603;
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  Matches
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EX BX SX B

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Gaps

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Indels

Length 20;

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Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
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                                                                                                                              The present sequence is a 3' PCR primer used in RT-PCR to amplify intron corresponding region from total P. pastoris mRNA, for isolation and characterisation of formaldehyde dehydrogenase 1 (FLD1) gene. FLD protects cells from toxic effects of formaldehyde and imparts formaldehyde resistance. It is therefore useful as a selectable marker in methylotrophic yeast. Expression cassettes comprising FLD regulatory sequences (promoter and terminator) can be used to control the expression of heterologous proteins in transformed methylotrophic yeast cells. Formaldehyde resistant host cells can be selected by transforming them with a vector comprising FLD gene operably linked to a heterologous promoter and growing them in the presence of formaldehyde.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAT00817 and AAT00818 are amplification primers for the stem, transmembrane and cytoplasmic domains of the 75 kD tumour necrosis factor
                                                                   useful as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tumour necrosis factor; TNF; erythropoletin receptor; EpoR; cytokine; tumour therapy; polymerase chain reaction; PCR; primer; amplify; ss.
                                                                                                                                                                                                                                                                                                                            Gaps
                                                              New formaldehyde dehydrogenase gene for methylotrophic yeast, us selection marker, also its promoter for regulated expression of heterologous polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New constructs for stimulating signalling in cells - comprising cytokine receptor cytoplasmic domain linked to a multimerising extracellular domain
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                                                                                                                                                                                                                                                                                                      Length 22;
                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                   Score 13.2; DB 21;
Pred. No. 1.4e+03;
0; Mismatches 3;
                                                                                                                                                                                                                                                                    Sequence 22 BP; 8 A; 5 C; 6 G; 3 T; 0 other;
(RESE ) RESEARCH CORP TECHNOLOGIES INC.
                                                                                                            Example 2; Page 46; 101pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 5; Page 16; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                           AAT00817 standard; DNA; 33 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5' 75 kD TNF receptor primer.
                                                                                                                                                                                                                                                                                                   Query Match 66.0%;
Best Local Similarity 83.3%;
Matches 15; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    95WO-US04221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94US-0224593
                                                                                                                                                                                                                                                                                                                                               3 tgatgcttaaaagcttac 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (TEXA ) UNIV TEXAS SYSTEM.
                                                                                                                                                                                                                                                                                                                                                           21 TGATGCTTACACGCTTTC 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Beutler BA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1995-358585/46.
                                           WPI; 2000-182118/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9526985-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-MAY-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-0CT-1995.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bazzoni FM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                       Cregg JM;
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The amplified sequence is used in a chimeric protein with the erythropoletria receptor (EDOR) sequence amplified by AATOOB13 and AATOOB14.

AATOOB14. A vector that expresses a chimeric receptor, such as the one created using the amplified sequence, can be used for the treatment of created using the amplified sequence, can be used for the treatment of tumours, and to provide a model for studying the formation of tumours. By altering the extracellular and transmembrane domains of the chimeric protein, the target sequence is changed. The formation of multimeric receptor complexes on the extracellular surface, when the chimeric protein is used, is sufficient for transmitting a positive signal to the cytoplasmic domain of the receptor. This occurs even in the absence of the cognate ligand, or in the presence of a ligand that has entirely different activities. The advantage with using the chimeric protein is the response of the cytokine (in this case TNF) can be induced
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This oligonucleotide was used as a primer for the PCR amplification of a cDNA clone (see AAX25358) coding for the T cell receptor (TCR) beta chain (see AAX05728) from the Al CD4+ T cell clone isolated from CBA/Ca mice. The Al clone recognises the minor histocompatibility antigen H-Y present in male, but absent in female, mice. The amplified product was used in the construction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         T cell receptor beta chain; mouse; transgenic animal; animal model; immunological tolerance; graft rejection; tissue grafting; neonatal intolerance; transplantation antigen; H-Y antigen; PCR; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 33;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 13.2; DB 16;
Pred. No. 1.4e+03;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mouse Al T cell receptor beta chain PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 33 BP; 8 A; 10 C; 6 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cobbold SP, Waldmann H, Zelenika D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 9; 41pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           66.0%;
83.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ouery Match
Best Local Similarity 83.39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   without side effects.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-255090/21.
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Identifying recombinantly an antimicrobial bioactive peptide used as a therapeutic agent involves transforming a host cell with expression vector with tightly regulable control region and measuring its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Escherichia coli; E. coli; randomised peptide library; identification; stabilised bioactive peptide; synthesis; intracellular selection; screening; lac operon; protease resistant; peptidase resistant; Rop protein; glutathione sulphotransferase; thioredoxin; infection; maltose binding protein; glutathione reductase; antimicrobial; antibacterial; PCR primer; ss.
                                                                                                                                                                                                                                                                                                         Gaps
             animal model comprising TCR alpha and beta chains. It provides a genetically modified non-human mammal having a population of CD4 positive T cells specific for one or a limited number of selected antigens, including at least transplantation antigen capable of rejecting a tissue transplant containing the transplantation antigen and if applicable, the other selected antigens. The animal has TCR genes which encode a TCR specific for the transplantation antigen. The animal is useful for studying immunological tolerance, especially the mechanisms of tolerance to, and the rejection of, tissue grafts, and in pregnancy. The animals are also useful for testing agents for biological activity in
    The invention relates to a transgenic
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                                                                                                                                                                                                                                                                        Length 33;
                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Plasmid pLAC11-recA construction PCR primer SEQ ID NO:17.
                                                                                                                                                                                                                                                                        promoting or reducing immunological tolerance.
                                                                                                                                                                                                                           Sequence 33 BP; 10 A; 5 C; 9 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   UYGE-) UNIV GEORGIA RES FOUND INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA56045 standard; DNA; 40 BP
                                                                                                                                                                                                                                                                        66.0%;
83.3%;
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                                                                                                                                                                                                                                                                                                                                     1 catgatgcttaaaagctt 18
                                                                                                                                                                                                                                                                                                                                                       05-SEP-2000 (first entry)
    of Al(M) transgenic mice.
                                                                                                                                                                                                                                                                                       Best Local Similarity 83.3
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-317972/27.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Escherichia coli.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-0CT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA56045;
                                                                                                                                                                                                                                                                          Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                   AAA56045,
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cell under conditions that repress expression of (P) and then inducing its expression which, if is inhibitory to host cell growth, is indicative of BP expression. An antimicrobial peptide from the present invention, which is stabilised, is used for treating a patient having a condition treatable with a peptide drug. The stabilised peptides are also used for inhibiting the growth of a microbe. The new antibacterial peptides are useful to treat various pathogenic bacteria such as staphylococci, Streptococci and Enterococci which are the primary causes of nosocomial infections. Over inhibitor peptides identified by the method can be medical treatments and therapies directed against microbial infection. Also, these novel inhibitor peptides can be used, in turn, to identify additional novel antibacterial peptides using a synthetic approach, and can also be used to elucidate potential new drug targets. The inhibitor peptide target which is inactivated is identified using reverse genetics by isolating mutants that are no longer inhibited
                                                                                                                                                                                                                                                                                       by the peptide. These mutants are then mapped in order to precisely determine the protein target that is inhibited. AAA56033 to AAA56106 and AAY90964 to AAY90999 are sequences used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Preparation of new human kernon acceptor co-repressor coding series and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes a human homologue of nuclear receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             coreceptor (HNRCR). The present sequence represents a PCR primer for human HNRCR, which is used in an example from the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 13.2; DB 21; Length 40; Pred. No. 1.4e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; HNRCR; nuclear receptor coreceptor; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (XINH-) XINHUANGPU FUDAN GENE ENG CO LTD SHANGHA.
                                                                                                                                                                                                                                                                                                                                                                                Sequence 40 BP; 12 A; 6 C; 5 G; 17 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP.
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83.3%;
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                                                                                                                                                                                                                                                                                                                                                 present invention.
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65.0%; Score 13; DB 21; Length 32;

Query Match

The present invention describes a method for identifying a bioactive peptide (BP) involving transforming a cell with an expression vector comprising a tightly regulatable control region operably linked to a nucleic acid sequence encoding a peptide (P), growing the transformed

Example 1; Page 36; 135pp; English.

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0; Gaps Best Local Similarity 100.0%; Pred. No. 1.8e+03; Matches .13; Conservative 0; Mismatches 0; Indels

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Search completed: October 2, 2001, 16:18:46 Job time: 15490 sec

22, Appli 3, Appli 3, Appli 15, Appli 30, Appli 20, Appl

Sequence Sequence Sequence

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Sequence 18, Application US/09377310B
Patent No. 6133031
GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION UNMBER: US/09/377,310B
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
LENGTH: 20
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APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Garden, William A.
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310B
CURRENT FILIAG DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SEQ ID NO 38
SEQ ID NO 38
LENGTH: 15
US-09-342-749-18

US-08-13711-24

US-08-133-011-22

US-08-133-011-22

US-08-122-730A-22

US-08-122-730A-22

US-08-132-730A-22

US-08-133-132-15

US-08-133-08-155-15

US-08-907-739-22

US-08-907-739-22

US-08-907-739-22

PCT-US93-08-364-22

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Matches 20;
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Sequence 106, App
Sequence 45, Appl
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Sequence 3, Appli
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Sequence 38, Sequence 38, Sequence 43, A
Sequence 11, S
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Sequence 39,
Sequence 10,
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Sequence 32,
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/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
                           GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-377-310-18
US-09-377-310-18
US-09-377-310-38
US-08-762-428A-10
US-08-569-959-32
US-08-569-959-32
US-08-958-835A-41
US-08-938-835A-41
US-08-938-835A-41
US-08-473-327-4
US-08-931-327-4
US-08-910-327-4
US-08-910-327-4
US-08-511-556-4
US-08-511-556-4
US-08-511-556-4
US-08-612-840A-9
US-08-612-840A-9
US-09-377-310-43
US-08-612-840A-9
US-08-612-840A-9
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US-08-617-958-9
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US-09-262-773-45
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                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Timothy F. Murphy TITLE OF INVENTION: Vaccine For Branhamella catarrhalis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM: MS-DOS/ Microsoft Windows 3.1 SOFTWARE: Wordperfect for Windows 5.1 SOFTWARE: Wordperfect for Windows 5.1 APPLICATION DATA: APPLICATION NUMBER: US/08/306,871 FILING DATE: 20-SEP-1994 PRIOR APPLICATION NUMBER: US 08/129,719 FILING DATE: September 29, 1993 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.4 Mb storage
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Pred. No. 4.6e+02;
); Mismatches 2;
                                                                                                                                                                                                                                                                                                              Score 12.8; DB 3;
Pred. No. 4.5e+02;
); Mismatches 2;
    REFERENCE/DOCKET NUMBER: 19603/1330
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STRAIN: 25240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 32, Application US/08306871
Patent No. 5712118
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 29 nucleotides
TYPE: nucleic acid
STRANDEDNESS: single-stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 35,300
REFERENCE/DOCKET NUMBER: 11
TELECOMMUNICATION INFORMATION: TELEPHONE: (716) 856-400
TELEFAX: (716) 849-0349
INFORMATION FOR SEQ ID NO: 32:
                         TELECOMMUNICATION INFORMATION:
TELEPHONE: 716-263-1636
TELEFAX: 716-263-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Hodgson, Russ, A
STREET: 1800 One M&T Plaza
CITY: Buffalo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 87.5%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: New York
COUNTRY: United States
ZIP: 14203-2391
                                                                    TELEFAX: 716-263-1600 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                              Query Match 64.0
Best Local Similarity 87.5
Matches 14; Conservative
                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 25 base pairs
TYPE: nucleic acid
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                                                                                                                                                                              STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nelson, M. Bud
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 52
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                       ; TOPOLOGY: linear; MOLECULE TYPE: CDNAUS-08-762-428A-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
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APPLICANT: Brett P. Monia
APPLICANT: Brett P. Monia
APPLICANT: Lex M. COWSETT
TILLE OF INVENTION: ANTISENSE MODULATION OF LIVER GLYCOGEN PHOSPHORYLASE EXPRESSION
FILE REPERENCE: RTS-0074
CURRENT APPLICATION NUMBER: US/09/357,071
CURRENT FILING DATE: 1999-07-19
SEQ ID NOS: 47
SEQ ID NOS: 47
LENGTH: 20
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71.0%; Score 14.2; DB 3; Length 20;
Best Local Similarity 84.2%; Pred. No. 83;
Matches 16; Conservative 0; Mismatches 3; Indels
                                                                                                         Ouery Match
75.0%; Score 15; DB 3; Length 15;
Best Local Similarity 100.0%; Pred. No. 31;
Matches 15; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: NIXON, HARGRAVE, DEVANS & DOYLE LLP STREET: Clinton Square, P.O. Box 1051
CITY: Rochester STATE: New York
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 14603
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/762,428A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: Antisense Oligonucleotide US-09-357-071-39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 4
US-08-762-428A-10
IS-08-762-428A-10
Sequence 10, Application US/08762428A
Fatent No. 6120993
GENERAL INFORMATION:
APPLICANT: Ye, Guo-jie
APPLICANT: Meister Alton
TITLE OF INVENTION: 5-OXOPROLINASE
NUMBER OF SEQUENCES: 10
CONRESPONDENCE ADDRESS:
ADDRESSEE: NIXON, HARGRAVE, DEVANS
             ; OTHER INFORMATION: antisense sequence US-09-377-310-38
                                                                                                                                                                                                                                                                                                                                                          Sequence 39, Application US/09357071 Patent No. 6043091
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 34,103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: TIMIAN, SUSAN J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 tttcaaccagatggtcatt 19
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                                                                                                                                                                                                                                                                                                              RESULT 3
US-09-357-071-39
FEATURE:
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Sequence 41. Application US/08938835A
Fatent No. 6060245
GENERAL INFORMATION:
APPLICANT: SORGE, Joseph A.
TITLE OF INVENTION: METHODS AND ADAPTORS FOR GENERATING
TITLE OF INVENTION: SPECIFIC NUCLEIC ACID POPULATIONS
NUMBER OF SEQUENCES: 69
CORRESPONDENCE ADDRESS:
ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
ADDRESSEE: Dunner, L.L.P.
STREET: 1300 I Street, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 12.6; DB 1; Length 35;
Pred. No. 5.9e+02;
0; Mismatches 4; Indels
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                                                                                                      COMPUTES: THE COMPUTED COMPUTES: THE COMPUTED COMPUTES: THE COMPUTED COMPUTES: THE COMPUTES: DOS SOFTWARE: FASTESQ Version 1.5 CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/469,005A FILING DATE: 05-JUN-1995
CLASSIFICATION: 536
PRIOR APPLICATION NUMBER: 08/188,426
FILING DATE: 24-JUL-1994
APPLICATION NUMBER: 08/096,908
FILING DATE: 26-JUL-1993
APPLICATION NUMBER: 07/917,716
FILING DATE: 24-JUL-1993
APPLICATION NUMBER: 34,688
REGISTRATION NUMBER: 34,688
REGERENCE/DOCKET NUMBER: 062482-0113
TELECOMMUNICATION: NUMBER: 062482-0113
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-639-7700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
ZIP: 2005-3315
COMPUTER READABLE FORM:
MEDUN TYRED: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      63.0%;
78.9%;
                                                                   MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 35 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 63.0
Best Local Similarity 78.9
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 202-639-7890
                                                   COMPUTER READABLE FORM:
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                           20004-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: line
MOLECULE TYPE: (
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; ANTI-SENSE: NO
; FRAGMENT TYPE:
; ORIGINAL SOURCE:
US-08-469-005A-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 8
US-08-938-835A-41/c
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                                                                                                                                Sequence 32, Application US/08569959
Patent No. 5725862
GENERAL INFORMATION:
APPLICANT: Timothy F. Murphy
TITLE OF INVENTON: Vaccine For Branhamella catarrhalis
NUMBER OF SEQUENCES: 52
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                         ADDRESSEE: Hodgson, Russ, Andrews, Woods & Goodyear
STREET: 1800 One M&T Plaza
CITY: Buffalo
                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 14203-2331

COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.4 Mb storage COMPUTER: IBM Compatible
COMPUTER: Wordperfible
OPERATING SYSTEM: MS-DOS/ Microsoft Windows 3.1
SOFTWARE: Wordperfect for Windows 5.1
CURREWY APPLICATION DATA:
APPLICATION NUMBER: US/08/569,959
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Patent No. 5665874
GENERAL INC. SECONDATION:
APPLICANT: KUHAJDA, FRANCIS P.
APPLICANT: PASTERNACK, GARY A.
TITLE OF INVENTION: CANCER RELATED ANTIGEN
UNMAREN OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: BAKER & BOTTS, L.L.P. STREET: 1299 Pennsylvania Avenue, N.W.-CITY: Washington STATE: DC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
PRIOR APPLICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 0.S. 08/129,719
FILING DATE: September 29, 1993
ATTORNEY/AGENT INFORMATION:
NAME: Nelson, M. Bud
REGISTRATION NUMBER: 35,300
REFERENCE/DOCKET NUMBER: 11520.0053
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPAX: (716) 849-0349
INFORMATION FOR SEQ ID NO: 32:
SEQUENCE CHARACTERISTICS:
TENGURLE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Branhamella catarrhalis
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EDNESS: single-stranded
GY: linear
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87.5%;
                                                                                                                                                                                                                                                                                                                                                             STATE: New York
COUNTRY: United States
ZIP: 14203-2391
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity 87.5
Matches 14; Conservative
10 TTCATCCAGATGATCA 25
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ORGANISM: Braun-
TW: 25240
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US-08-469-005A-7/c
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                                                                                                               US-08-569-959-32
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US-08-569-959-32
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                                                                                                                                Length 30;
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                                                                                                                                                                          1; Indels
                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: WEIL ET AL.
TITLE OF INVENTION: habe
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
ADDRESSEE: OARELLA, BYINE, BAIN, GILFILLIAN,
ADDRESSEE: CECCHI, STEWART & OLSTEIN
STREET: 6 BECKER FARM ROAD
CITY: ROSELAND
STATE: NEW JERSEY
COUNTRY: USA
                                                                                                                                  Score 12.4; DB 1;
Pred. No. 7.4e+02;
); Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5 INCH DISKETTE
COMPUTER: 18M PS/2
OPERATING SYSTEM: MS-DOS
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/783,266
FILING DATE: 15-JAN-1997
CLASSIFICATION 1435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/463,975
FILING DATE: 10no 5,1995
APPLICATION NUMBER: PCT/US94/12058
FILING DATE: 21 OCT 1994
ATTONNEY, AGENT THORMATION:
MANAT: FRORMATON:
                                                                                                                                                                                                                                                                                                         RESULT 10
US-08783-266-4
; Sequence 4, Application US/08783266
; Patent No. 5747312
                                      ; TOPOLOGY: LINEAR
; MOLECULE TYPE: Oligonucleotide
US-08-463-975-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; MOLECULE TYPE: Oligonucleotide US-08-783-266-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-994-1700
                                                                                                                                  62.0%;
92.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: FERRARO, GREGORY D. REGISTRATION NUMBER: 36,1 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 201-994-1744
INFORMATION FOR SEQ ID NO: 4:
                                                                                                                              Query Match
Best Local Similarity 92.9
Matches 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             : 30 BASE PAIRS
NUCLEIC ACID
TYPE: NUCLEIC ACID
STRANDEDNESS: SINGLE
                                                                                                                                                                                                                                  9 TTTCATCCAGATGG 22
                                                                                                                                                                                                              1 tttcaaccagatgg 14
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                                    LINEAR
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Matches 13; Conser
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Pred. No. 7.2e+02;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 4, Application US/08463975
Patent No. 5618717
GENERAL INFORMATION:
APPLICANT: WEI, ET AL.
TITLE OF INVENTION: habH
NUMBER OF SEQUENCES:
CORPESSEE: CARELLA, BYRNE, BAIN, GILFILLAN,
ADDRESSEE: CRECHI, STEWART & OLSTEIN
STREET: 6 BECKER FARM ROAD
CITY: ROSELAND
STATE: NEW JERSEY
COUNTRY: USA
ZIE: 07068
COMPUTER READABLE FORM:
MEDIUM TYPE: 35,5 INCH DISKETTE
        FILING DATE: 26-SEPT-1997

FILING DATE: 26-SEPT-1997

CLASSIFICATION:
PRIOR APPLICATION:
APPLICATION NUMBER: US 08/775,993

FILING DATE: 03-JAN-1997

PRIOR PAPLICATION NUMBER: US 08/779,335

FILING DATE: 06-JAN-1997

ATTORNEY-AGENT INFORMATION:
NAME: BATKET, NFORMATION:
NAME: BATKET, NFORMATION:
TELEPHONE: 202-408-4000

TELEPHONE: 202-408-4000

TELEPHONE: 26-408-4000

INFORMATION FOR SEQ ID NO: 41:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYPE: nucleic acid
STANDEDRESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA (Genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM: MS-DOS
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NAMER: US/08/463,975
FILING DATE: June 5, 1995
CLASSIPECATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMER: PCT/US94/12058
FILING DATE: 21 OCT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FERRARO, GREGORY D.
REGISTRATION NUMBER: 36,134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 36,134
REFERENCE/DOCKET NUMBER: 325800-331
TELECOMMUNICATION INFORMATION:
TELEBRONE: 201-994-1700
TELEFAX: 201-994-1744
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     62.0%;
92.9%;
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INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
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Best Local Similarity 92.9
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PS/2
OPERATING SYSTEM: P
                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-938-835A-41
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Score 12.2; DB 1; Length 20;
Pred. No. 8.9e+02;
0; Mismatches 3; Indels
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Patent No. 5783390
GENERAL INFORMATION:
APPLICANT: Agoulnik, A. APPLICANT: Kent, Marijo G. TITLE OF INVENTION: ATTLE OF INVENTION: ATTLE OF INVENTION: BATTERY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: WILLIAM STATE: WILLIAM STATE: WILLIAM STATE: WILLIAM STATE: USA ZIP: 53717-1914
COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPPRATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PATENTIN Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/472,416
FILLING DATE: US/03/1995
CLASSIFICATION: 435
                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
GURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/531,556
                                                                                                                                                                  CLASSIFCATION: 435
ATTORNEY_AGENT INFORMATION:
NAME: Sara, Charles S.
RECISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 34506.034CIP
TELEPHONE: 608-831-2100
TELEFAX: 608-831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 94
CORRESPONDENCE ADDRESS:
ADDRESSER: Dewitt Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
CITY: Madison
STATE: WI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 3450
REFERENCE/DOCKET NUMBER: 3450
TELECOMMUNICATION INFORMATION:
TELEPHONE: 608-831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     61.0%;
                        Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 TCCATCCAGCTGGTCAT 17
                                                                                                                                                                                                                                                                                                                            TELEFAX: 608-831-2106
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 ttcaaccagatggtcat 18
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 61.0
Best Local Similarity 82.4
Matches 14; Conservative
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                        single
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-531-556-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-472-416-4
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APPLICANT: Agoulnik, Alexander I
APPLICANT: Rent First, Marijo
APPLICANT: Meallem, Ariege
TITLE OF INVENTION: MALE INFERTILITY Y-DELETION DETECTION
TITLE OF INVENTION: BATTERY
NUMBER OF SEQUENCES: 124
CORRESPONDENCE ADDRESS:
ADDRESSEE: DeWitt Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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                                                   GENERAL INFORMATION:
APPLICANT: WEI, ET AL.
TITLE OF INVENTION: hABH
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
ADDRESSEE: CARELLA, BYRNE, BAIN, GILFILLAN,
ADDRESSEE: CECCHI, STEWART & OLSTEIN
STREET: 6 BECKER FARM ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/463,975
FILING DATE: June 5, 1995
FILING DATE: 21 OCT 1994
ATJOHNEY/AGENT INFORMATION:
NAME: FERRARO, GREGORY D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: FERRARO, GREGORY D.
REGISTRATION NUMBER: 36,134
REFERENCE/DOCKET NUMBER: 325800-331
TELECOMUNICATION INFORMATION:
TELEPHONE: 201-994-1700
TELEFAX: 201-994-1744
                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/09/023,327
                                                                                                                                                                                                                                                                                                   2IP: 07068

COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 INCH DISKETTE
COMPUTER: IBM PS/2
               Sequence 4, Application US/09023327 Patent No. 5929225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 4, Application US/08531556 Patent No. 5776682
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; MOLECULE TYPE: Oligonucleotide US-09-023-327-4
                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: MS-DOS
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 201-994-1744 INFORMATION FOR SEQ ID NO: 4:
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LENGTH: 30 BASE PAIRS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 62.0
Best Local Similarity 92.9
Matches 13; Conservative
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TYPE: NUCLEIC ACID
STRANDEDNESS: SINGLE
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                                                                                                                                                                                                                                      CITY: ROSELAND
STATE: NEW JERSEY
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: LINEAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                53717-1914
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: W. COUNTRY:
US-09-023-327-4
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US-08-859-998-1026
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                                                                                                                                          Gaps
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US-09-101-886B-30
; Sequence 30, Application US/09101886B
; Sequence 30, Application US/09101886B
; Batent No. 6197507
; GENERAL INFORMATION:
    APPLICANT: BERG, THOMAS
; APPLICANT: TOLLERSRUD, OLE K
; APPLICANT: TOLLERSRUD, OLE K
; APPLICANT: NILSSEN, OIVIND
; TITLE OF INVENTION: GENETIC TEST FOR ALPHA-WANNOSIDOSIS
; NUMBER OF SEQUENCES: 104
; CORRESPONDENCE ADDRESS:
    ADDRESSEE: BADRESS:
    ADDRESSEE: S55 13TH STREET, NW SUITE 701E
    CITY: WASHINGTON
    STATE: DC
COUNTRY: USA
                                                                                               Score 12.2; DB 1; Length 20;
Pred. No. 8.9e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 12.2; DB 4; Length 23;
Pred. No. 9e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 20004
COMPUTER READABLE FORM:
MEDIUM TYPE: RIOPY disk
COMPUTER: ISM COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: 29-JANUARY-1998
CLING DATE: 29-JANUARY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /desc = "oligonucleotide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRICE APPLICATION DATA:
APPLICATION NUMBER: PCT/GB97/00109
FILING DATE: 12-JAN-11997
ATTORNEY/AGENT INFORMATION:
NAME: ERNST, BARBARA G
REGISTRATION NUMBER: 30,377
REFERENCE/DOCKET NUMBER: 1181-240
TELEPHONE: 202-783-6040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "oligonucl
; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
US-08-472-416-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61.0%;
82.4%;
                                                                                             Ouery Match 61.0%;
Best Local Similarity 82.4%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 202-783-6031
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
LENGTH: 23 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                             2 ttcaaccagatggtcat 18
                                                                                                                                                                                                      1 TCCATCCAGCTGGTCAT 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3 tcaaccagatggtcatt 19
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TELEPHONE: 202-783-6031
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Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; ANTI-SENSE:
US-09-101-886B-30
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Sequence 1026, Application US/08859998
Patent No. 5994076
GENERAL INFORMATION:
APPLICANT: Chenchik, Alex
APPLICANT: Jokhadze, George
APPLICANT: Jokhadze, George
APPLICANT: Bibilashvilli, Robert
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 1375
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 12.2; DB 2;
Pred. No. 9.3e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                      COUNTY 15 05
ZIP: 94025
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: 1BM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FASCESO for Windows Version 2.0
CURRENT APPLICATION DATE: APPLICATION NUMBER: US/08/859,998
FILING DATE: 21-MAY-1997
CLASSIFICATION 1435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Oligonucleotide primer US-08-859-998-1026
                                                                                                                                                                                                        ADDRESSEE: Fish & Richardson, P.C. STREET: 2200 Sand Hill Road, Suite 100 CITY: Menlo Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09096/002001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2, 2001, 16:03:53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 09096/
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-322-5070
TELEFAX: 415-854-0875
INFORMATION FOR SEQ ID NO: 1026:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61.0%;
82.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 28 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Search completed: October
Job time: 14597 sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity Matches 14; Conserv
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Tyrosine-Kinase sy Tyrosine-Kinase sy Tyrosine-Kinase sy PCR primer used in Probe used in prep Probe used to isol Micro gene random RBP-7 biallelic ma Primer #17 used in ER-specific probe. PCR primer used to PCR primer for amplify

Chitinase specific Chitinase degenera SPT6 homologue for Loci-specific prim Human NBC PCR prim Polymorphic fragme

Human homologue of Human AlkB homolog

Human NBC PCR prim Human wolframin in

PCR primer PDZK5.4 Oestrogen receptor Dog genomic marker Human focal adhesi

Human beta-1,3-gal Human beta-1,3-gal B.catarrhalis CD e Polymorphic fragme

Rat 5-oxoprolinase

Neisseria species Synthetic plasmid

score:

Sequence: Title: Perfect s

nucleic

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Scoring table:

Searched:

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Database

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New antisense compounds for inhibiting focal adhesion kinase expression, especially useful for inhibiting retinal neovascularization, or for diagnosing and treating e.g. colon cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; focal adhesion kinase; FAK; signal transduction; cancer; embryonic development disorder; angiogenic disorder; wound healing; antisense; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human focal adhesion kinase antisense sequence #16.
                                                                                                                                                                                                                                                                                                                                                                                                                                        ALIGNMENTS
                                                          AAA53029
AAQ91896
AAA04221
                                                                                                                                                       AAX03260
AAT78192
                       AAX52031
AAV28114
                                                                                              AAQ73036
AAQ73037
                                                                                                                                                                                                        AAN92881
AAV40115
                                                                                                                                                                                                                                          AAT68318
AAX88777
                                                                                                                                                                                                                                                                                          AAT72839
AAT68289
                                                                                                                                                                                                                                                                                                                                         AAV72184
AAZ88895
                                                                                                                                  AAT88208
                                                                                                                                              AAX15118
                                                                                                                                                                                 AAZ87025
                                                                                                                                                                                            AAC90273
                                                                                                                                                                                                                                AAT29698
                                                                                                                                                                                                                                                                   AAC66377
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                                                                                                                                                                                                                                                                                                                                                                                         AAA66803
                                                                                                                                                                                                                                                                                                                                                                                                    AAC65575
                                               AAA53025
                                                                                                                       AAA29977
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP.
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 AAC65550 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
  (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Monia BP, Gaarde WA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-006141/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US6133031-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-OCT-2000
AAC65550;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAC65550
  RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Maize polymorphic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human focal adhesi
Human focal adhesi
PCR primer PDZK5.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human liver glycog
                                                                              Search time 876.95 Seconds (without alignments)
14.320 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Description
                                                                                                                                                                                                                                                                                                                                                           | Sings | Sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /SIDS8/gcgdata/geneseq/geneseqn/NA2000.DAT:*/SIDS8/gcgdata/geneseq/geneseqn/NA2001.DAT:*
        4.5
Compugen Ltd.
                                                                                                                                                                                                                                      Potal number of hits satisfying chosen parameters:
                                                                                                                                                                                                                 730101 seqs, 313950809 residues
                                                                                  ٠.
           GenCore version
Copyright (c) 1993 - 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SUMMARIES
                                                                                2001, 16:18:45
                                                                                                                                                                                                                                                                                                    Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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AAC86411
AAA14039
AAA14039
AAV51245
AAV51246
AAV47956
AAV47957
AAV12243
AAV51243
AAV51243
                                                         - nucleic search, using sw model
                                                                                                                                                                              IDENTITY_NUC
Gapop 10.0 , Gapext 1.0
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20
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Match Length
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Maximum DB seq length: 50
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Score

Result Š 14.4 113.8 113.8 113.8 113.8 113.4 113.4

5 6 7 7 8 8 9 11

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Bartel PL,
                                                                                                                                                                                                  Synthetic
                                                                                                 AAX86411;
                                                       RESULT
AAX86411/
ID AAX8
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                   The present invention describes a number of phosphorothioate antisense sequences to the human focal adhesion kinase (FAK) protein. This protein is involved in integrin-mediated signal transduction, and is implicated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes a number of phosphorothioate antisense sequences to the human focal adhesion kinase (FAK) protein. This protein is involved in integrin-mediated signal transduction, and is implicated in cancer, particularly colon, breast and oral tumours, embryonic development disorders, anglogenic disorders and wound healing. The antisense sequences, including the one shown here, can be used in the treatment of all of these.
                                                   in cancer, particularly colon, breast and oral tumours, embryonic development disorders, anglogenic disorders and wound healing. The antisense sequences, including the one shown here, can be used in the treatment of all of these.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense compounds for inhibiting focal adhesion kinase expression, especially useful for inhibiting retinal neovascularization, or for diagnosing and treating e.g. colon cancer
                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                      Human; focal adhesion kinase; FAK; signal transduction; cancer; embryonic development disorder; anglogenic disorder; wound healing; antisense; phosphorothloate; ss.
                                                                                                                                                                ;
0
                                                                                                                                          Length 20;
                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                 Human focal adhesion kinase antisense seguence #36.
                                                                                                                                         DB 22;
0.22;
                                                                                                          Sequence 20 BP; 5 A; 5 C; 3 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15 BP; 5 A; 4 C; 3 G; 3 T; 0 other;
                                                                                                                                                                Mismatches
                                                                                                                                         Score 20;
Pred. No. (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 2; Column 25; 30pp; English
 Claim 3; Column 23; 30pp; English
                                                                                                                                                               ö
                                                                                                                                         100.0%;
                                                                                                                                                                                                1 tttcaaccagatggtcattc 20
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                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                     Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaarde WA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-006141/01.
                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-AUG-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                      US6133031-A.
                                                                                                                                                                                                                                                                                                           12-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Monia BP,
                                                                                                                                          Query Match
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MASC1 protein. The PCR templates were derived from tumour cell lines, and the amplicons were tested for mutations. The MMSC1 protein is a cand the amplicons were tested for mutations. The MMSC1 protein is a ward interacting protein which is involved in tumour suppression activity in the MMAC1 pathway. MMSC1, antigenic fragments or fusion continuity in the MMAC1 pathway. MMSC1, antigenic fragments or fusion continuity in the MMAC1 genemates for antibody production. Primers derived from MMSC1 gene used as immunogens for antibody production of MMSC1 can be used for identification of MMSC1 gene an alteration in am MMSC1 gene is indicative of a predisposition to cancer. A seminine cancer. A somatic mutation in an MMSC1 gene is indicative of a predisposition to cancer. A somatic mutation in an MMSC1 gene is indicative that the tissue is cancerous. Analysis of MMAC1 and MMSC1 or predisposition to cancer. Sometic mutations can be used for detection of alterations in MMAC1 cancer. Wild-type MMSC1 or a homologue can be used to supply wild-type MMSC1 gene function (or a substantially similar concertion) to a cell, which has lost the gene function due to a MMSC1 gene mutation. The gene suppresses neoplastic growth of the cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                             Human; MMSC1 protein; MMAC1 interacting protein; tumour suppression;
MMAC1 pathway; immunogen; cancer; cell neoplastic growth; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MMSC1, an MMAC1 (tumour suppressor) interacting protein and related
                                                                                                                                                                                                             PCR primer PDZK5.5Q used to amplify DNA encoding MMSC1 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            identifying drug candidates useful in treating cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 14.4; DB 20;
Pred. No. 2e+02;
); Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         8 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 5; Page 50; 107pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              72.0%;
93.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99WO-US00995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98US-0071861
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (MYRI-) MYRIAD GENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 38 BP; 11 A; 10 C;
AAX86411 standard; DNA; 38
                                                                                                                                     29-SEP-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 72.0
Best Local Similarity 93.8
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polynucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9936566-A1
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RESULT

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Gaps

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0; Indels

75.0%; Score 15; DB 22; 100.0%; Pred. No. 83; iive 0; Mismatches 0,

Conservative

Query Match Best Local Similarity Matches 15; Conserv

Length 15;

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Polymorphic marker; allele-specific; primer; probe; amplification; hybridisation; plant; hybrid certification; genetic contribution; progeny; back-cross; hybrid; ancestry; maize; ss.
                                                                                                                                                                                              Maize polymorphic marker S02G2/G5-2 DNA
                  RESULT 5
AAV51245/c
ID AAV51245 standard; DNA; 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (AFFY-) AFFYMETRIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                         W09824796-A1
                                                                                                                                                                 11-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                       11-JUN-1998
                                                                                                                                                                                                                                                                                                                                variation
                                                                                                                                    AAV51245;
                                                                                                                                                                                                                                                                                       Zea mays.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequences AAA14008-A14047 represent phosphorothioate antisense oligonucleotides targetted to the human liver glycogen phosphorylase coligonucleotides targetted to the human liver glycogen phosphorylase sequence of interpression. The antisense coligonucleotides were designed to target different regions of human liver glycogen phosphorylase RNA, and were analysed for their effect on liver glycogen phosphorylase is one of three glycogen phosphorylase is encoded by three different genes. Liver glycogen phosphorylase is encoded by the PYGL gene, which is located on chromosome 14. Liver glycogen phosphorylase (also known as 1,4-alpha-D-glucan:orthophosphate cappropriate glycogen phosphorylase (also known as 1,4-alpha-D-glucan:orthophosphate cappropriate in the liver to glucose-1-phosphate via the cleavage of the alpha-1,4-glycosidic bonds. It therefore plays a critical role in carbohydrate metabolism and blood glucose homeostasis. Inhibition of liver glycogen phosphorylase and chere's in diabetic patients, particularly those with type II (non insulin-dependent) diabetes. The antisense olygonicototides collonuctootides of the invention are useful for diagnosis, prevention and treatment of conditions associated with liver glycogen phosphorylase expression, or those which may benefit from inhibition of liver glycogen phosphorylase
                                                                                                                                   1,4-alpha-D'glucan:orthophosphate alpha-D-glucosyltransferase; HGLP:
glycogenolysis, carbohydrate metabolism; blood glucose homeostasis;
expression inhibition; hypoglycaemic; type II diabetes;
non insulin-dependent; antisense; phosphorothioate; ss.
                                                                                       Human liver glycogen phosphorylase antisense oligo, SEQ ID NO:39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              diagnosis and treatment of diseases associated with
                                                                                                                     Liver glycogen phosphorylase; PYGL gene; human; chromosome 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antisense compounds particularly oligonucleotides useful for
                                                                                                                                                                                                                                                                      /*tag= a
/note= "Phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 5 A; 5 C; 4 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            expression of liver glycogen phosphorylase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   such as type II diabetes
                                                                                                                                                                                                                                       Location/Qualifiers
1..20
/*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 3; Column 40; 33pp; English.
AAA14039 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                            99US-0357071
                                                                                                                                                                                                                                                                                                                                                                                                        99US-0357071
                                                           18-JUL-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                      (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Monia BP, Cowsert LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2000-270346/23.
                                                                                                                                                                                                                                        Key
modified_base
                                                                                                                                                                                                                                                                                                                                                                            19-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                        19-JUL-1999;
                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                              28-MAR-2000.
                                                                                                                                                                                                                                                                                                                US6043091-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20
                            AAA14039;
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/note= "polymorphism"

97WO-US21782 97US-0813507 96US-0032069

/*tag= a /replace= "c"

Location/Qualifiers

BP

(first entry)

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                                                                                                                                                                                                              This DNA sequence is a region of a Zea mays genome which contains a polymorphic marker. This sequence can be used in the construction of allele-specific primers and probes for amplification or hybridisation. e.g. to determine common or disparate ancestry between 2 or more plants, i.g. monitor the genetic contribution of an ancestral plant, to trace the progeny of proprietary plants, in certification of a hybrid plant to to defentify the progeny of a back-crossed plant with an ancestral plant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                          Brassica species allele-specific oligonucleotide probes and primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ó
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
Sapolsky RJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 13.8; DB 19;
Pred. No. 4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 41 BP; 8 A; 8 C; 13 G; 12 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
  Murigneux A,
                                                                                                                                                                 Claim 1; Page 49; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAV51246 standard; DNA; 41 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               69.0%;
88.2%;
                                                                                                                      useful for plant breeding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4 caaccagatggtcattc 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22 CCACGAGATGGTCATTC
Lemieux B,
                                             WPI; 1998-333252/29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Landry BS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAV51246;
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ID AAV512
XX
AC AAV512
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Gaps

; 0

Query Match 71.0%; Score 14.2; DB 21; Length 20; Best Local Similarity 84.2%; Pred. No. 2.38+02; Matches 16; Conservative 0; Mismatches 3; Indels (

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The present invention describes a nucleic acid segment comprising at least 10 contiguous nucleotides from a vegetal sequence including a polymorphic site which is a single nucleotide polymorphism (SNP), or the complement of the segment. Also described are: (1) an allele-specific oligonucleotides hybridising to segment, or their complements, and (2) a method of analysing nucleic acids from a subject, by determining if a base is occupying any one (or a set) of polymorphic sites in 261 sequences derived from six maize lines (see AAW47701 to AAW47961). The segments are useful in fingerprint analysis in plants to determine which polymorphisms are present, which strain a plant belongs to and to distinguish between strains. The polymorphisms may correlate with phenotypic traits (e.g. plant growth rate or crop yield), and the segments are useful to determine the presence/absence of specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polymorphisms correlating with the existence/absence of particular traits. The segments are also useful in marker assisted back-cross techniques to select plants with a higher percentage of recurrent parent in a back-cross population. Segments incorporate SNRs which occur more frequently than other polymorphism types and are therefore more likely to be located close to genetic loci of interest; different forms of characterised SNRs are also often easier to detect than other
                                                                                                                                                                                                                                                                                                                              e.g. to determine polymorphisms in plants, determine strain in plant breeding and to correlate polymorphisms with phenotypic traits
                                                                                                                                                                                                                                                                                                            Vegetal sequences including single nucleotide polymorphism - useful, e.g. to determine polymorphisms in plants, determine strain in plant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Maize; marker; probe; PCR primer; polymorphism; vegetal sequence; polymorphic site; corn; graminae species; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Maize polymorphic site oligonucleotide marker UMC76-G2/G5-2B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 13.8; DB 19;
Pred. No. 4.2e+02;
); Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 41 BP; 8 A; 8 C; 12 G; 12 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                              Claim 2; Page 15; 32pp; English.
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ID AAV47957 standard; DNA; 41 BP.
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88.2%;
                                                                                      97WO-EP07134
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Matches 15; Conservative
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                                                                                                                                                                             (BIOC-) BIOCEM SA.
WO9830717-A2
                                                                                      02-DEC-1997;
                                                                                                                               02-DEC-1996;
                                                                                                                                                                                                                        Murigneux A;
                                         16-JUL-1998.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This DNA sequence is a region of a Zea mays genome which contains a polymorphic marker. This sequence can be used in the construction of allele-specific primers and probes for amplification or hybridisation, e.g. to determine common or disparate ancestry between 2 or more plants, to monitor the genetic contribution of an ancestral plant, to trace the progeny of proprietary plants, in certification of a hybrid plant or to identify the progeny of a back-crossed plant with an ancestral plant.
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                                                                                                      Polymorphic marker; allele-specific; primer; probe; amplification; hybridisation; plant; hybrid certification; genetic contribution; progeny; back-cross; hybrid; ancestry; maize; ss.
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88.2%; Pred. No. 4.2e+02;
.ive 0; Mismatches 2;
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                                                             Maize polymorphic marker S02G2/G5-2B DNA.
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/note= "polymorphism"
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96US-0032069
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Matches 15; Conservative
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variation
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WO9830717-A2

Synthetic.

AAV47956/C ID AAV479

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Zea sp.

16-JUL-1998.

Length 41; Indels

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This DNA sequence is a region of a Zea mays genome which contains a polymorphic marker. This sequence can be used in the construction of allele-specific primers and probes for amplification or hybridisation. e.g. to determine common or disparate ancestry between 2 or more plants, to contribution of an ancestral plant, to trace the progeny of proprietary plants, in certification of a hybrid plant or to identify the progeny of a back-crossed plant with an ancestral plant.
                                                                                                                                                                                                                                                                                                                          Brassica species allele-specific oligonucleotide probes and primers
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                                                                                                                                                                                                                                          Sapolsky RJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 41 BP; 8 A; 8 C; 12 G; 13 T; 0 other;
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/replace= "t"
/note= "polymorphism"
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96US-0032069.
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                                                                                                                                                                                              (AFFY-) AFFYMETRIX INC.
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                                                                                                                           07-MAR-1997;
02-DEC-1996;
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WO9824796-A1
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                                                                                                                                                                                                                                   Vegetal sequences including single nucleotide polymorphism - useful, e.g. to determine polymorphisms in plants, determine strain in plant breeding and to correlate polymorphisms with phenotypic traits
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                  97WO-EP07134
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Best Local Similarity
Matches 15; Conserv
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                  02-DEC-1997;
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                                                                                                                                                Murigneux A;
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AAV51243/c ID AAV512 RESULT

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AAV47955;
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                                                                                                    This DNA sequence is a region of a Zea mays genome which contains a polymorphic marker. This sequence can be used in the construction of allele-specific primers and probes for amplification or hybridisation, e.g. to determine common or disparate ancestry between 2 or more plants, to monitor the genetic contribution of an ancestral plant, to trace the progeny of proprietary plants, in certification of a hybrid plant or to identify the progeny of a back-crossed plant with an ancestral plant.
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                                               Brassica species allele-specific oligonucleotide probes and
                                                                                                                                                                                                                                                          Indels
 Sapolsky RJ;
                                                                                                                                                                                                                                  Score 13.4; DB 19;
Pred. No. 6.8e+02;
0; Mismatches 1;
                                                                                                                                                                                                 Sequence 41 BP; 8 A; 9 C; 11 G; 13 T; 0 other;
 Murigneux A,
                                                                                 Claim 1; Page 49; 65pp; English
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                                                                                                                                                                                                                                   67.0%;
93.3%;
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                                                           useful for plant breeding
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                                                                                                                                                                                                                                Query Match 67.0
Best Local Similarity 93.3
Matches 14; Conservative
                                                                                                                                                                                                                                                                                6 accagatggtcattc 20
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 Lemieux B,
                       WPI; 1998-333252/29
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 Landry BS,
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The present invention describes a nucleic acid segment comprising at least 10 contiguous nucleotides from a vegetal seguence including a polymorphic site which is a single nucleotide polymorphism (SNP), or the complement of the segment. Also described are: (1) an allele-specific oligonucleotides hybridising to segment, or their complements, and (2) a method of analysing nucleic acids from a subject, by determining if a base is occupying any one (or a set) of polymorphic sites in 261 sequences derived from six maize lines (see AAV47701 to AAV47961). The segments are useful in fingerprint analysis in plants to determine which polymorphisms are present, which strain a plant belongs to and to distinguish between strains. The polymorphisms may correlate with
are useful in fingerprint analysis in plants to determine which
                                                                                                                                                                                                                                                       techniques to select plants with a higher percentage of recurrent parent in a back-cross population. Segments incorporate SNPs which occur more frequently than other polymorphism types and are therefore more likely to be located close to genetic loci of interest; different forms of characterised SNPs are also often easier to detect than other
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                          polymorphisms are present, which strain a plant belongs to and to distinguish between strains. The polymorphisms may correlate with phenotypic traits (e.g. plant growth rate or crop yield), and the segments are useful to determine the presence/absence of specific polymorphisms correlating with the existence/absence of particular traits. The segments are also useful in marker assisted back-cross
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Maize; marker; probe; PCR primer; polymorphism; vegetal sequence; polymorphic site; corn; graminae species; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Maize polymorphic site oligonucleotide marker UMC76-G2/G5-1B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 13.4; DB 19;
Pred. No. 6.8e+02;
); Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 41 BP; 8 A; 8 C; 11 G; 13 T; 1 other;
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AAV47955 standard; DNA; 41 BP.
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93.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                 polymorphism types
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WO9914318-A1
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                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                  AAX52031;
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                                                                                                                                                                                                                                        RESULT 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ53015 to AAZ54536, AAZ54577 to AAZ54615, and AAY74253 to AAY75941 represent novel Neisseria meningitis and N. gonorrheae polynucleotides and polypeptides. AAZ54537 to AAZ54576 and AAZ54616 to AAZ55473 represent PCR primers used in the exemplification of the present invention. The
phenotypic traits (e.g. plant growth rate or crop yield), and the segments are useful to determine the presence/absence of specific polymorphisms correlating with the existence/absence of particular traits. The segments are also useful in marker assisted back-cross techniques to select plants with a higher percentage of recurrent parent in a back-cross population. Segments incorporate SNPs which occur more frequently than other polymorphism types and are therefore more likely to be located close to genetic loci of interest; different forms of characterised SNPs are also often easier to detect than other
                                                                                                                                                                                                                                                                                                                                                         Neisseria meningitidis; Neisseria gonorrheae; antigen; vaccine; antigenic; diagnosis; immunogenic; infection; meningitis; septicaemia; antibacterial; gene therapy; PCR primer; ss.
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                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel Neisserial polypeptides predicted to be useful antiqens for
                                                                                                                                                                         ö
                                                                                                                                                  67.0%; Score 13.4; DB 19; Length 41; 93.3%; Pred. No. 6.8e+02; ive 0; Mismatches 1; Indels 0
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Scalato E, S
                                                                                                                    Sequence 41 BP; 8 A; 9 C; 10 G; 13 T; 1 other;
                                                                                                                                                                                                                                                                                                                                     Neisseria species ORF cloning PCR primer #699.
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Ratti G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 16; Page 160; 1453pp; English.
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, Pizza M, Rappuoli R,
                                                                                                                                                                                                                                                                      AAZ55314 standard; DNA; 33 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98US-0094869.
98US-0098994.
98US-0099062.
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98US-0103794.
98US-0103796.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99US-0121528
                                                                                                                                                                                                                                                                                                               21-MAR-2000 (first entry)
                                                                                                                                                  Query Match 67.0
Best Local Similarity 93.3
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         vaccines and diagnostics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GENO-) INST GENOMIC RES
                                                                                                                                                                                            6 accagatggtcattc 20
                                                                                                                                                                                                        Petersen J, Pizza M,
Tettelin H, Venter JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-062150/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CHIR ) CHIRON CORP.
                                                                                                 polymorphism types.
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                                                                                                                                                                                                                                                                                                                                                                                                                Neisseria sp.
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09-OCT-1998;
25-FEB-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fraser C,
                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
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construct a synthetic DNA plasmid sequence synlus4, to demonstrate the method of the invention. Within the synlus4 sequence are included the sequences of lux A, lux B, the A and B components of the Vibrio fisherii luciferase sequence, positions of pUC19 including the origin of replication and replication stability sequences, and the promoter and coding sequence for the Ranamycin/neomycin phosphotransferase. The specification describes a method for the synthesis of replication competent double-stranded polynucleotides. The method comprises generating a first set of oligonucleotides corresponding to the plus strand and a second set corresponding to the minus strand and a second set corresponding to the minus strand and a nocoding sequences involved in a biochemical pathway. In particular, they can be used to produce polynucleotides encoding enzymes, they can be used to produce polynucleotides encoding enzymes, they can be used to produce polynucleotides encoding enzymes, they can be used to produce polynucleotides encoding enzymes, and tolase, thiose-phosphate isomerase, phosphoglycerate mutase, phosphoglycerate mutase,
polypeptides, the polynucleotides, antibodies and compositions of the invention can be used as vaccines, as diagnostic reagents, and as immunogenic compositions. The polypeptides can be used in the manufacture of medicaments for treating or preventing infection due to Neisserial bacteria (e.g. meningitis and septicaemia), to detect the presence of Neisseria bacteria, or to raise antibodies. They may also be used to screen for agonists or antagonists, which may themselves have use as antibacterial agents. The polynucleotides of the invention may also be used in gene therapy protocols.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthesis of replication competent double-stranded polynucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA plasmid; lux A; lux B; Vibrio fisherii; luciferase; promoter; tn9 kanamycin/neomycin phosphotransferase; DNA synthesis; replication competent double-stranded polynucleotide; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 33;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 13.2; DB 21;
Pred. No. 8.3e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 33 BP; 11 A; 10 C; 4 G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 4; Fig 5A; 135pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 83.3
Matches 15; Conservative
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Search completed: October 2, 2001, 16:18:45 Job time: 15489 sec

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enolase or pyruvate kinase. They can also be used for the preparation of viral particles, artificial genomes and artificial genetic systems
                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This PCR primer was used in the amplification of 5-oxoprolinase cDNA (see also AAV28111-12) from rat kidney cDNA. The primer incorporates an Ndel site at the start codon of the 5-oxoprolinase. CDNA. The PCR product was used in the construction of expression plasmids for 5-oxoprolinase. Mammalian 5-oxoprolinase (m50P) catalyses the ATP dependent cleavage of 5-oxoproline to L-glutamate in the metabolism of glutathione. Deficiency of the enzyme is associated with 5-oxoprolinuria. The invention is directed toward isolated nucleic acid molecules encoding m50P. Expression vectors and host cells are provided, as well as methods of increasing (by gene therapy) or decreasing (e.g. using antisense or ribozyme molecules) the expression of m50P in host cells.
                                                                                                                                                                                                                                                                                                                                                                                                    5-oxoprolinase; rat; glutathione; 5-oxoprolinuria; gene therapy; diagnosis; PCR; primer; ss.
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                                                                                            Length 50;
                                                                                                                            3; Indels
                                                                                           Score 13.2; DB 20;
Pred. No. 8.9e+02;
0; Mismatches 3;
                                              Sequence 50 BP; 21 A; 9 C; 8 G; 12 T; 0 other;
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                                                                                                                                                                                                                                                                           BP.
                                                                                                                                                                                                                                                                                                                                                                    Rat 5-oxoprolinase PCR primer.
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                                                                                            Query Match 66.0%;
Best Local Similarity 83.3%;
Matches 15; Conservative
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                                                                                                                                                                                                                                                                       AAV28114 standard; DNA; 25
                                                                                                                                                           1 tttcaaccagatggtcat 18
                                                                                                                                                                            14 tttctaccagatgataat 31
                                                                                                                                                                                                                                                                                                                                      26-OCT-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       treating 5-oxoprolinuria
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Rattus sp.
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Score 12.8; DB 19; Length 25; Pred. No. 1.3e+03; 0; Mismatches 2; Indels (

Query Match
Best Local Similarity 87.5%;
Matches 14; Conservative

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Sequence:

Title:

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Sequence 17, Application US/09377310B

Sequence 17, Application US/09377310B

Patent No. 6133031

GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310B
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 17
FERMAND: APPLICATION OF 12.0
   Sedinence Sedine
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US-09-037-143-16
US-09-049-651-16
US-08-260-174-16
US-08-59-1095
US-08-58-6719-2
US-08-58-6719-2
US-08-58-6719-2
US-07-866-560-2
US-07-866-560-2
US-08-077-673-2
US-08-077-673-2
US-08-077-673-2
US-08-077-673-2
US-08-077-673-2
US-08-077-673-2
US-08-077-673-2
US-08-077-673-19
US-08-08-08-110-19
US-08-08-110-19
US-08-08-110-19
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US-08-08-110-19
US-08-08-18-19
US-08-08-18-19
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Fatent No. 5765209
GENERAL INFORMATION: GENERAL INFORMATION: FUNCTIONAL DOMAINS OF DIA
TITLE OF INVENTION: FUNCTIONAL DOMAINS OF DNA
NUMBER OF SEQUENCES: 67
CORRESPONDENCE ADDRESS:
ADDRESSE: Annold, White & Durkee
STREET: 31 No. 5763209th Clark Street, Sui
STATE: 1111nois
CITY: Chicago
STATE: 1111nois
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                          ALIGNMENTS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; FEATURE:
; OTHER INFORMATION: antisense sequence
US-09-377-310-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;
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MEDIUM TYPE: Floppy disk
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20; Conservative
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US-09-377-310-17
 Query Match
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                                                                                                                              Search time 417.38 Seconds
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/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/PcTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/PcTUS_COMB.seq:*
                     GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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US-08-46-52
US-08-44-52
US-08-686-968C-20
US-08-684-422-16
US-08-614-45-16
US-09-082-614-16
US-08-647-3518-3
US-08-647-3518-3
US-08-647-3518-3
US-08-647-348-3
US-08-637-448-3
US-08-637-448-3
US-08-637-448-19
US-08-639-670-3
US-08-639-670-3
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US-08-639-670-3
US-08-639-670-3
US-08-639-670-3
US-08-639-190-3
US-08-639-190-3
US-08-430-190-3
US-08-430-133-764-5
PCT-US94-07091-1
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US-08-124-981A-18
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US-09-037-192-16
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Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 1.0
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Pred. score and is

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Sequence 37, Application US/09377310B
Patent No. 6133031
GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
TITLE OF INVENTION: Expression
TITLE OF INVENTION: Expression
TITLE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310B
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 37
LENGTH: 15
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                                                                                                        DB 1; Length 40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.8; DB 4; Length 32;
Pred. No. 1.1e+02;
); Mismatches 2; Indels
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APPLICANT: Cochran, Mark D.
TITLE OF INVENTION: Recombinant Swinepox Virus
FILE REFERENCE: 39119-H/JML
CURRENT APPLICATION NUMBER: US/08/686,968C
CURRENT FILING DATE: 1996-07-25
NUMBER OF SEQ ID NOS: 231
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 20
LENGTH: 32
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. 82;
                                                                                                    76.0%; Score 15.2; D
85.0%; Pred. No. 72;
Live 0; Mismatches
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Mismatches
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100.0%; Pred. No.
tive 0; Mismatc
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; Sequence 20, Application US/08686968C
; Patent No. 6221361
; GENERAL INFORMATION:
                       ; MOLECULE TYPE: DNA (genomic) US-08-466-344-52
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88.9%;
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                                                                                                      Query Match 76.0%
Best Local Similarity 85.0%
Matches 17; Conservative
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Best Local Similarity 88.9°
Matches 16; Conservative
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      TOPOLOGY: linear
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Best Local Similarity
Matches 15; Conserv
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TITLE OF INVENTION: METHODS AND MATERIALS RELATING TO THE
TITLE OF INVENTION: FUNCTIONAL DOMAINS OF DNA BINDING PROTEINS
NUMBER OF SEQUENCES: 67
CORRESPONDENCE ADDRESS:
ADDRESSEE: Aroll, White & Durkee
STREET: 321 No. 5773583th Clark Street, Suite 800
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 40;
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APPLICATION NUMBER: US/08/466,344
FILING DATE: 06-JUN-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/040,548
FILING DATE: 08-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: COUGHLIN, Daniel F.
REGISTRATION NUMBER: 36,111
REFERENCE/DOCKET NUMBER: 36,111
REFERENCE/DOCKET NUMBER: arcd067.
TELECOMMUNICATION INFORMATION:
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATEMIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/040,548
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                   CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: COUGHLIN, Daniel F.
REGISTRATION NUMBER: 36,111
REFERENCE/DOCKET NUMBER: arcd0
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 744-0090
INFORMATION FOR SEQ ID NO: 52:
SEQUENCE CHARACTERISTICS:
LENGTH: 40 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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Patent No. 5773583
                                                                                                                                                                                                                                                                                                                                                                                                       ) MOLECULE TYPE: DNA (genomic) US-08-040-548-52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (312) 744-0090
TELEFAX: (312) 245-4961
INFORMATION FOR SEO ID NO: 52:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity 85.03
Matches 17; Conservative
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TYPE: nucleic acid
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STRANDEDNESS:
                                                                                                    FILING DATE:
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TITLE OF INVENTION: Rapid Detection of Antibiotic Resistance TITLE OF INVENTION: in Mycobacterium Tuberculosis NUMBER OF SEQUENCES: 66
ADDRESSPONDENCE ADDRESS:
ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 18;
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                                                               STATE: MEDITING CONTRY: USA

ZIP: 20005-3315
COMPUTER: ELOPPY disk
COMPUTER: IBM PC COMPA: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/313,185
FILING DATE: 12-OCT-1994
CLASSIFICATION: 435
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COMPUTER: IBM PC COMPATIBLE
OPERATES: SYSTEM: PC-DOS/MS-DOS
SOFTWARE: 'Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 2;
                                                                                                                                                                                                                                                                                                                                                        ATTORNEY AGENT INFORMATION:
NAME: MAYORS, Kenneth J.
REGISTRATION NUMBER: 25,146
REFERENCE, DOCKET NUMBER: 0336.0068-00000
TELECOMMUNICATION INFORMATION:
TELEBRAN: (202) 408-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 13.4; DB 2;
Pred. No. 4.6e+02;
0; Mismatches 1
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APPLICATION NUMBER: US/09/082,614A
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 16, Application US/09082614A Patent No. 6124098 GENERAL INFORMATION:
                      STREET: 1300 I Street, N.W. CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Cole, Stewart
APPLICANT: Young, Douglas
APPLICANT: Zhang, Ying
APPLICANT: Honore, Nadine
APPLICANT: Telenti, Amalio
APPLICANT: Telenti, Amalio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 1300 I Street, N.W. CITY: Washington STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
US-08-313-185-16
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93.3%;
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ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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16 CTCGCTGCGGGTGGA 2
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  Dunner
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Best Local Similarity
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US-09-082-614A-16/c
ADDRESSEE:
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TITLE OF INVENTION: in Mycobacterium Tuberculosis
                                                                                        Score 14.2; DB 2; Length 20;
Pred. No. 2e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Finnegan, Henderson, Farabow, Garrett &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/628,422
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04020/094001
                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Fish & Richardson P.C. STREET: 225 Franklin Street CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 16, Application US/08313185; Patent No. 5851763; GENERAL INFORMATION: APPLICANT: Heym, Beate APPLICANT: Cole, Stewart APPLICANT: Young, Douglas APPLICANT: Zhang, Ying APPLICANT: Honore, Nadine APPLICANT: Telenti, Amalio APPLICANT: Bodmer, Thomas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Fasse, J. Peter
REGISTRATION NUMBER: 32,983
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ó
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
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84.2%;
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                      15 attettegetgatggtgg 32
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Best Local Similarity 84.2
Matches 16; Conservative
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TELEX: 200154
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; MOLECULE TYPE: DNA
US-08-628-422-16
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Gaps

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; TOPOLOGY: linear
US-08-861-450-2
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US-09-392-580-43/c
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US-08-647-351B-3
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APPLICANT: Br inker, Peter
APPLICANT: Minas, Wolfgang
APPLICANT: Malla, Pauli
APPLICANT: Malla, Pauli
APPLICANT: Mallay, James E.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR
TITLE OF INVENTION: INCREASING PRODUCTION OF ERITHROMYCIN
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                              Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                    1; Indels
                                                                                                                                                                                                                                                                                                                                                                                              Score 13.4; DB 3;
Pred. No. 4.6e+02;
0; Mismatches 1;
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/313,185
FILING DATE: 12-OCT-1994
ATTONNEY/AGENT INFORMATION:

NAME: Meyers, Kenneth J.

REGISTRATION NUMBER: 25,146
REFERENCE/DOCKET NUMBER: 25,146
REFERENCE/DOCKET NUMBER: 02356.0068-0000
TELECOMMUNICATION INFORMATION:
TELEPRAX: (202) 408-400
TELEPRAX: (202) 408-400
TELEPRAX: (202) 408-400
TELEPRAX: LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDBUSS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CUBRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/861,450
FILING DATE: 22-MAY-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9092-0004-999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     E: Pennie & Edmonds, LLP
1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 2, Application US/08861450 Patent No. 5908764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 9092
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-5556
TELEFAX: 650-493-5556
TELEX: 66141 PRINIE
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                ; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
US-09-082-614A-16
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93.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 93.3
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Abrams, Samuel B
REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
ZIP: 10036-2811
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5 ctcgctgctggtgga 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
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STATE:
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Sequence 43, Application US/09392580
Patent No. 608173
GENERAL INFORMATION:
APPLICANT: C. Frank Bennett
APPLICANT: Lex M. Cowsert
APPLICANT: Lex M. Cowsert
ITILE OF INVENTION: ANTISENSE MODULATION OF X-LINKED INHIBITOR OF APOPTOSIS EXPRES
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                                              Indels
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                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Sheldon & Mak
STREEF: 225 S. Lake Avenue, 9th Floor
CITY: Pasadena
STATE: California
STATE: California
COMPUTER READBLE FORM:
MEDIUM TYRE: Diskette, 3.50 inch, 1.44 Mb storage
COMPUTER: IBM COMpatible
OPERATING SYSTEM: Windows 95
SOFTWARE: WordPerfect for Windows version 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/647,351B
FILING DATE: MAY 9, 1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: De Leon, Ricardo
APPLICANT: Rochelle, Paul
TITLE OF INVENTION: Cryptosporidium Detection Method
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
    Score 13.4; DB 2;
Pred. No. 4.8e+02;
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); Mismatches
                                              0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Other nucleic acid primer sequence
                                                                                                                                                                                                                              ; Sequence 3, Application US/08647351B
; Patent No. 5770368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Farah, David A.
REGISTRATION NUMBER: 38,134
REFERENCE/DOCKET NUMBER: 113
TELECOMMUNICATION INFORMATION:
  67.0%;
93.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  64.0%;
87.5%;
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TELEFAX: (626) 795-6321
INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                            14; Conservative
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Matches 14; Conservative
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                                                                                    6 tcgctgctggtggaa 20
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                                                                                                                           26 TAGCTGCTGGTGGAA 12
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; MOLECULE TYPE:
; DESCRIPTION:
US-08-647-351B-3
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Score 12.6; DB 1; Length 30;
Pred. No. 1.2e+03;
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                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRICEATION: 435
PRICEATION 108TA:
APPLICATION NUMBER: US 08/295,670
FILING DATE: 08-SEP-1994
APPLICATION NUMBER: PCT/JP94/00039
FILING DATE: 13-JAN-1994
PRICE APPLICATION NUMBER: 19 5-4069
FILING DATE: 13-JAN-1993
ATORNEY/AGENT INFORMATION:
RAGISTRATION NO. 5681717man F.
REGISTRATION NUMBER: 24,618
REGISTRATION NUMBER: 24,618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/633,485
FILING DATE: 17-APR-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; TOPOLOGY: linear
; MOLECULE TYPE: other nucleic acid
US-08-633-485-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDLUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ELECOMMUNICATION INFORMATION
                   63.0%;
78.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (703) 413-3000
TELEFAX: (703) 413-2220
TELES: 248855 OPAT INCORMATION FOR SEQ ID NO: 3:
                                                                                                       2 ttcctcgctgctggtggaa 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 ttcctcgctgctggtggaa 20
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                                                                                                                                   1 TTCATCGCTGCTGTCGCA 19
                                   Best Local Similarity 78.9
Matches 15; Conservative
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LENGTH: 30 base pairs
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Best Local Similarity
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US-08-633-485-3
                   Query Match
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ADDRESSEE: P.C.
                                                                                                                                                                                                                                                                                                                 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 1755.5. Jefferson Davis Highway, Suite 400 CITY: Arlington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 3, Application US/08295670
Patent No. 5547864
GENERAL INFORMATION:
APPLICANT: TAMASAKI, HISASHI
APPLICANT: TSUCHIYA, MAKOTO
APPLICANT: MWA, KIYOSHI
APPLICANT: MWA, KIYOSHI
APPLICANT: MAMAHARA, YOSHIO
TITLE OF INVENTION: THE NOVEL CELL SURFACE PROTEIN
NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                   Score 12.6; DB 3;
Pred. No. 1.1e+03;
0; Mismatches 4;
                                                                                                                                                              FEATURE:
; OTHER INFORMATION: Antisense Oligonucleotide
US-09-392-580-43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 19-5-4069
FILING DATE: 13-JAN-1993
ATTORNEY/AGENT INFORMATION:
NAME: Oblon, No. 5547864man F.
REGISTRATION NUMBER: 24,618
REFERENCE/ADCKET NUMBER: 10-697-0 PCT
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 413-3000
                 CURRENT APPLICATION NUMBER: US/09/392,580
CURRENT FILING DATE: 1999-09-09
NUMBER OF SEQ ID NOS: 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/295,670
FILING DATE: 08-5EP-1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/JP94/00039
FILING DATE: 13-JAN-1994
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TOPOLOGY: linear; MOLECULE TYPE: other nucleic acid us-08-295-670-3
                                                                                                                        TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                   Query Match 63.0%;
Best Local Similarity 78.9%;
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                        1 attectegetgetggtgga 19
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TELEX: 248855 OPAT UR
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                              19 ATTGTTCCATGCTGGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 30 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
FILE REFERENCE: RTS-0072
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: Virginia COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22202
                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 12
US-08-295-670-3
                                                                               SEQ ID NO 43
LENGTH: 20
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Gaps ö ö Sequence 3, Application US/08633485
Patent No. 5681717
GENERAL INFORMATION:
APPLICANT: RAWASARI, HISASHI
APPLICANT: TSUCHITA, MAKOTO
APPLICANT: KAWAHARA, YOSHI
APPLICANT: KAWAHARA, YOSHI
TITLE OF INVENTION: THE NOVEL CELL SURFACE PROTEIN
NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,
ADDRESSEE: P.C.
STREET: 1755 S. Jefferson Davis Highway, Suite 400
CITY: Arlington
STATE: Virginia 63.0%; Score 12.6; DB 1; Length 30; 78.9%; Pred. No. 1.2e+03; ive 0; Mismatches 4; Indels Indels

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Search completed: October 2, 2001, 16:03:52 Job time: 14596 sec
                       OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: WORDPERFECT 5.1
    IBM PC COMPATIBLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 61.0%;
Best Local Similarity 82.4%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 attectegetgetggtg 17
                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TOPOLOGY:
US-08-430-225A-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Dong, Jin-Tang; Barrett,
APPLICANT: J. Carl; Lamb, Patricia W.; Isaacs, John T.
TILE OF INVENTION: DIAGNOSTIC METHODS AND
TITLE OF INVENTION: GENE THERAPY USING REAGENTS DERIVED FROM THE
TITLE OF INVENTION: HUMAN METASTASIS SUPPRESSOR GENE KAII
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORGAN & FINNEGAN, L.L.P.
STREET: 345 PARK AVENUE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ő
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61.0%; Score 12.2; DB 1; Length 20; 82.4%; Pred. No. 1.7e+03; 1ve 0; Mismatches 3; Indels
                                  Sequence 4, Application US/08202042

Petent No. 5686072

GENERAL INFORMATION:

APPLICANT: Jonathan W. Uhr

APPLICANT: Bilen S. Vitetta

TITLE OF INVENTION: EPITOPE-SPECIFIC MONOCLONAL

TITLE OF INVENTION: ANTIBODIES AND IMMUNOTOXINS

TITLE OF INVENTION: AND USES THEREOF

NUMBER OF SEQUENCES: 6

CORRESPONDENCE ADDRESS:

ADDRESSEE: Arnold, White & Durkee

STREET: P.O. Box 4433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/202,042
FILING DATE: Submitted herewith
FILING DATE: Submitted herewith
FILING DATE: Submitted herewith
ATTORNEY/AGENT INFORMATION:
NAME: David L. Parker
REFERENCE/DOCKET NUMBER: UTSD:379/PAR
TELECOMMUNICATION INFORMATION:
TELEPHONE: (512) 474-7577
INFORMATION FOR SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy Disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 13, Application US/08430225A Patent No. 6204000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4 ATTTCTCGCTCGTGGTG 20
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Best Local Similarity 82.4
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                              ZIP: 77210
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy I
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                                                                                                                                                                                                                                                                                                                                       Texas: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USA
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US-08-430-225A-13
                                                                                                                                                                                                                                                                                                              CITY: Hou
STATE: TE
COUNTRY:
RESULT 14
US-08-202-042-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ò
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ZIP: 10154 COMPUTER READABLE FORM: MEDIUM TYPE: FLOPPY DISK

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Gaps
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Pred. No. 1.7e+03;
); Mismatches 3; Indels
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/430,225A
FILING DATE: 28-APR-1995
                                                                                                                                                   2026-4172
                                                         CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: RICHARD W. BORK
REGISTRATION NUMBER: 2026
REFERENCE/DOCKET NUMBER: 2026
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 758-4800
TELERX: (212) 751-6849
INFORMATION FOR SEQ ID NO: 13: SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
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3 65.0 23 21 AAC87185 3 65.0 23 21 AAC88395 3 65.0 23 21 AAC89319 8 64.0 19 21 AAC8953 8 64.0 21 18 AAR8992 6 63.0 20 AAX97166	12.6 63.0 20 21 AAA64943 12.6 63.0 30 15 AAQ70365 12.4 62.0 18 14 AAQ37914 12.4 62.0 18 17 AAQ37914	12.4 62.0 21 21 AAAA8928 12.4 62.0 29 21 AAF056016 12.4 62.0 31 20 AAX02733 12.4 62.0 31 21 AAX45459 12.4 62.0 31 21 AAX45459	27 12.2 61.0 20 18 AAT93237 28 12.2 61.0 20 20 AAS04613 29 12.2 61.0 21 17 AAT40022	12.2 61.0 21 19 AAV57633 12.2 61.0 24 15 AAQ56130 12.2 61.0 29 14 AAQ56130	33 12.2 61.0 30 AAX16994 34 12.2 61.0 31 16 AAQ76298 35 12.2 61.0 31 18 AAX62574	12.2 61.0 31 21 AAA/9051 12.2 61.0 38 19 AAN81428 12.2 61.0 38 21 AAA/57997	12.2 61.0 38 21 AAA0/288 12.2 61.0 41 21 AAC73359 12.2 61.0 42 16 AA086682 12.2 61.0 42 21 AA000913 12.2 61.0 42 21 AAA29268	45 12.2 61.0 42 22 AAF31700 ALIGNMENTS	ESULT 1 AC65549 D AAC65549 standard; DNA; 20 BP. X	C AAC65549; X T 12-FEB-2001 (first entry)	Human focal adhesion Human, focal adhesic embryonic developmen	Homo sapiens. US6133031-A.	7	F 19-AUG-1999; 99US-0377310.	Hd SISI (-SISI)	Monia	WPI; 2001-006141/01.	T New Antisense compounds for inhibiting local adnession Kinase T expression, especially useful for inhibiting retinal T neovascularization, or for diagnosing and treating e.g. colon cancer -
12 1 13 1 14 1 15 12. 16 12.	Seconds c 18 12. 5) 20 12. 1 updates/sec 21 12.	23 12. 24 12. 25 12. 25 12.	27 12. 28 12. 29 12.	30 12. 31 12. 32 12.	c 34 12.	12.	2 40 12. C 41 12. C 42 12. C 44 12.	c 45 12.	RESULT 1 AAC65549 ID AAC65549 XX	AC A XX XX DT 1	XX XX XX XX XX XX XX	ve a XX XX XX printed, XX XX XX	17	19	XX (ISIS-)	AX PI Monia XX	used to XX	
GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd. OM nucleic - nucleic search, using sw model	Run on: October 2, 2001, 16:18:44 ; Search time 876.95 (without alignment: 14.320 Million cell	Title: US-09-757-100B-17 Perfect score: 20 Sequence: 1 attcctcgctgctggtggaa 20	Scoring table: IDENTITY_NUC Gapop 10.0 , Gapext 1.0	Searched: 730101 seqs, 313950809 residues Total number of hits satisfying chosen parameters: 8549	eq length: 0 eq length: 50	Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries	Database: N_Geneseq_0601:* 1: /SIDSB/gcgdata/geneseq/geneseqn/NA1980.D 2: /SIDSB/gcgdata/geneseq/geneseqn/NA1981.D 3: /SIDSB/gcgdata/geneseq/geneseqn/NA1982.D	4: /SIDSB/gcgdata/geneseq/geneseqn/NA1983.DAT:* 5: /SIDSB/gcgdata/geneseq/geneseqn/NA1984.DAT:* 6: /SIDSB/gcgdata/geneseq/geneseqn/NA1985.DAT:* 7: /SIDSB/gcgdata/geneseq/geneseqn/NA1986.DAT:* 8: /SIDSB/gcgdata/geneseq/geneseqn/NA1986.DAT:*	. SIDSB/gcgdata/geneseq/geneseqn/NA1988.D 10: /SIDSB/gcgdata/geneseq/geneseqn/NA1989. 11: /SIDSB/gcgdata/geneseq/geneseqn/NA1990. 12: /SIDSB/gcgdata/geneseq/geneseqn/NA1991. 13: /SIDSB/gcgdata/geneseq/geneseqn/NA1991.	14: /SIDS8/gcgdata/geneseq/geneseqn/NA1993.15: /SIDS8/gcgdata/geneseq/geneseqn/NA1994.16: /SIDS8/gcgdata/geneseq/geneseqn/NA1995.	17: /SIDSB/gcgdata/geneseq/geneseqn/NA1996.DAT:* 18: /SIDSB/gcgdata/geneseq/geneseqn/NA1996.DAT:* 19: /SIDSB/gcgdata/geneseq/geneseqn/NA1998.DAT:* 20: /SIDSB/gcgdata/geneseq/geneseqn/NA1999.DAT:* 21: /SIDSB/gcgdata/geneseq/geneseqn/NA1999.DAT:* 23: /SIDSB/gcgdata/geneseq/geneseqn/NA1999.DAT:*	Pred. No. is the number of results predicted by chance to h score greater than or equal to the score of the result bein and is derived by analysis of the total score distribution.		Result Query No. Score Match Length DB ID	20 100.0 20 22 15 75.0 15 22	5 14.8 74.0 35 19 4 14.8 74.0 35 21 5 14.2 71.0 20 18	14 70.0 18 14 14 70.0 18 21	0 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6

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Swinepox virus; SPV; recombinant; vaccine; immunisation; diagnosis; pseudorabies virus; feline immunodeficiency virus; FIV; heartworm; Dirofilaria immitis; PCR primer; ss.
                                                                                                                                                                            IBRV gI gene PCR primer SEQ ID NO:20 from WO9804684 Example 41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Recombinant swine pox virus – useful in vaccine for immunising animal against swine pox virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 41; Page 180; 473pp; English.
                                                                                             AAV26137 standard; DNA; 32 BP.
                                                                                                                                                 24-JUL-1998 (first entry)
                3 tectegetgetggtg 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1998-130677/12.
                                                                                                                                                                                                                                                                                                                                                                                                (SYTR ) SYNTRO CORP.
                                                                                                                                                                                                                                                                                                                                           25-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                     25-JUL-1996;
                                                                                                                                                                                                                                                                                       WO9804684-A1
                                                                                                                                                                                                                                                                                                                                                                                                                           Cochran MD,
                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                          AAV26137;
                                                                   RESULT
AAV26137
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                         The present invention describes a number of phosphorothioate antisense sequences to the human focal adhesion kinase (FAK) protein. This protein is involved in integrin-mediated signal transduction, and is implicated in cancer, particularly colon, breast and oral tumours, embryonic development disorders, angiogenic disorders and wound healing. The antisense sequences, including the one shown here, can be used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a number of phosphorothioate antisense sequences to the human focal adhesion kinase (FAK) protein. This protein is involved in integrin-mediated signal transduction, and is implicated in cancer, particularly colon, breast and oral tumours, embryonic development disorders, anglogenic disorders and wound healing. The antisense sequences, including the one shown here, can be used in the treatment of all of these.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New antisense compounds for inhibiting focal adhesion kinase expression, especially useful for inhibiting retinal neovascularization, or for diagnosing and treating e.g. colon cancer
                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; focal adhesion kinase; FAK; signal transduction; cancer;
embryonic development disorder; angiogenic disorder; wound healing;
antisense; phosphorothioate; ss.
                                                                                                                                                                                                       ö
                                                                                                                                                                           100.0%; Score 20; DB 22; Length 20; 100.0%; Pred. No. 1.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     75.0%; Score 15; DB 22; Length 15; 100.0%; Pred. No. 2.9e+02; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                            Human focal adhesion kinase antisense sequence #35.
                                                                                                                                     Sequence 20 BP; 3 A; 5 C; 6 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 15 BP; 0 A; 5 C; 5 G; 5 T; 0 other;
                                                                                                                                                                                                     Mismatches
Example 2; Column 23; 30pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; Column 25; 30pp; English,
                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                             AAC65569 standard; DNA; 15 BP
                                                                                                                                                                                                                                1 attcctcgctgctggtggaa 20
                                                                                                                                                                                                                                              1 attectegetgetggtggaa 20
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                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                          treatment of all of these.
                                                                                                                                                                                       Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaarde WA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-006141/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                12-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US6133031-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-0CT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Monia BP,
                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                      AAC65569;
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AAC65569
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97WO-US12212. 9608-0686968

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The present sequence represents a PCR primer used in an example from the present invention. The present invention specifically describes recombinant swinepox virus (SPV) comprising a foreign DNA (I) inserted into a SPV genome which is capable of being expressed in a host cell into which the virus is introduced, where (I) is inserted into: (a) an ECORI site within a region corresponding to a 3.2 kb subfragment of the HindIII k fragment which contains both a HindIII and an ECORI site, of the SPV genome, and optionally (b) an AccI site within a region corresponding to a 3.6 kb HindIII to BajlII subfragment of the HindIII M cragment. The recombinant SPV can be used in a vaccine for immunising an animal against SPV. The invention also provides a method for testing a swine to determine whether the swine has been vaccinated with the accine, particularly containing S-SPV-008, or is infected with a naturally occurring wild-type pseudorables virus. Also (I) inserted into immunodeficiency virus (FIV) env and agg genes and Dirofilaria immitis psy and 22kd are useful to detect feline immunodeficiency caused by FIV and to detect heartworm caused by D. immitis respectively.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 14.8; DB 19;
Pred. No. 3.9e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 32 BP; 4 A; 7 C; 10 G; 11 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 74.0%;
88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 attectegetgetggtgg 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 16; Conserv
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RESULT AAA70370

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Gaps

; 0

Conservative

AAA70370;

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This antisense oligonucleotide is specific to a portion of the BZLFI gene. Antisense oligonucleotides specific to any of the different strains of Epstein-Barr virus (EBV) such as the BZLFI, BRLFI, BMLFI, BW THOSE ANTISCHES OLIGONUCLEOTIGES AND ANTISCHES OLIGONUCLEOTIGES AND ANTISCHES ANTIS
                              Epstein-Barr virus; antisense oligonucleotide; EBV; treatment; tumour;
BZLF1; BRLF1; BMLF1; BLLF1; BNLF1; BYRF1; BARF1; BZLF2; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     catalase-peroxidase; isonicotinic acid hydrazide; isoniazid; INH; antibiotic; susceptibility; sensitive; resistant; katG; polymerase chain reaction; Single strand conformation polymorphism; PCR-SSCP analysis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Anti:sense oligo:nucleotide(s) that inhibit Epstein-Barr virus function - used to treat or prevent EB virus infections, including
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         71.0%; Score 14.2; DB 18; Length 20; 84.2%; Pred. No. 7.1e+02; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Primer pair #6 Oligo2 to amplify M.tuberculosis katG gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 5 A; 7 C; 4 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                efficiency and reduce their needed dosage.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 5; Page 54; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ61472 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                          97WO-US06104.
                                                                                                                                                                                                                                                                                                                                                                                    96US-0628422.
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                                                                                                                                                                                                                                                                                                                                                                                                                                           (UYMA-) UNIV MASSACHUSETTS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1997-512403/47.
                                                                                                                   Synthetic.
Epstein-Barr virus.
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                                                                                                                                                                                                                                                                                                                       04-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                    05-APR-1996;
                                                                                                                                                                                                      WO9737669-A1
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                                                                                                                                                                                                                                                                16-0CT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ouery Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Key
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Op
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
0
                                                                                                                                                                                                                              Mucociliary dysfunction; mucus; sputum; human;
chronic obstructive lung disease; chronic bronchitis; CB; Bronchiectasis;
BE; asthma; cystic fibrosis; CF; bacterial infection; placental bikunin;
Kunitz-type serine protease inhibitor; chronic sinusitis; glue ear; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mucociliary dysfunction is the inability of ciliated epithelium to clear mocus and sputum in lung airways. Mucociliary dysfunction is a serious complication of chronic obstructive lung diseases such as Chronic bronchitis (CB). Bronchiectasis (BE), asthma and Cystic Fibrosis (CF). In addition, patients suffering from mucociliary dysfunction are susceptible to secondary bacterial infections. A partial coding sequence for human placental bikuin has been isolated (see AAA70371). Placental bikuin is a Kunitz-type serine protease inhibitor protein, which can stimulate the rate of mucociliary clearance of mucus and sputum in lung airways. Therefore, placental bikuin protein may be used for treating ung diseases such as CF, CB, BE, and chronic sinusitis and glue ear which are caused by retention and accumulation of mucus. The present sequence is an oligonucleotide used to clone the partial human placental bikunin coding sequence of AAA70371.
                                                                                                                                                                       3' oligonucleotide # 1 used for cloning human placental bikunin cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Stimulating mucociliary clearance rate of mucus and sputum in lung airways for treating lung diseases such as cystic fibrosis and bronchitis involves administering a Kunitz-type serine protease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 14.8; DB 21;
Pred. No. 3.9e+02;
); Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 35 BP; 7 A; 9 C; 14 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Poll CT, Newton BB, Taylor WJA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 6; Page 52; 173pp; English.
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88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98US-0218913.
99US-0441966.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99WO-GB04381
AAA70370 standard; DNA; 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tectegetgetggtggaa 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-APR-1998 (first entry)
                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 74.0
Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2000-452127/39.
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                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22-DEC-1998;
17-NOV-1999;
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                                                                                                                02-FEB-2001
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inhibitor

Hall R,

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AAT92645/c

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23-FEB-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                              Shen B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sugars
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Loca
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17
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AAQ88809
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   οy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eleven pairs of primers were synthesised and used to generate PCR products of around 280bp, that covered the complete katG gene and some of the flanking sequences. With these primers, a panel of INH-resistant strains of M tuberculosis was examined using PCR-SSCP analysis to localise the katG mutations responsible for isoniazid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rapid detection of antibiotic resistance in Mycobacteria - esp. 1soniazid, rifampicin or streptomycin resistance in tuberculosis by detecting mutation in katG, rpoB or rpsL genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR primer used to amplify bleomycin (BLM) gene cluster ORF34.
                   /*tag= a
/note= "letter O printed at this site in the
specification"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                         Telenti A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 14; DB 14;
Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18 BP; 5 A; 6 C; 5 G; 1 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                  Cole S, Heym B, Honore N, Zhang Y;
                                                                                                                                                                                                                                                                                                                                                (UYBE-) UNIV BERNE.
(UYPA-) UNIV CURIE PARIS VI P & M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 2; Page 29; 97pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        70.0%;
93.3%;
                                                                                                                                                                                                                    920S-0875940
                                                                                                                                                                                                                                                                                          (ASSI-) ASSISTANCE PUBLIQUE.
(INSP ) INST PASTEUR.
(MEDI-) MEDICAL RES COUNCIL.
                                                                                                                                                                93WO-EP01063
                                                                                                                                                                                                      92FR-0011098
                                                                                                                                                                                                                                                          93FR-0004545
                                                                                                                                                                                                                                                                                                                                 MEDICAL RES COUNCIL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-JAN-2000; 2000WO-US00445.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA58527 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 70.0
Best Local Similarity 93.3
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Streptomyces verticillus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1993-368812/46.
misc_difference 8
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                                                                                                                                                                30-APR-1993;
                                                                                                                                                                                                                    30-APR-1992;
14-AUG-1992;
                                                                                                                                                                                                                                                          16-APR-1993;
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                                                                                       W09322454-A
                                                                                                                            11-NOV-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    resistance
                                                                                                                                                                                                                                                                                                                                                                                                       Bodmer T,
                                                                                                                                                                                                                                                                                                                                                                                                                         Young D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA58527;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR primers AAA58474-A58541 were used to amplify open reading frames (ORFs) 8 to 41 of the BLM (Bleomycin) gene cluster. The proteins encoded by the gene cluster are useful for producing peptides and/or polyketide metabolites, especially bleomycin or bleomycin analogues. They are also useful for chemically modifying biological molecules to produce branched methyl groups, and for coupling amino acids and fatty acids. They may be reacted with an apo-carrier protein and coenzyme A to produce a holo-carrier protein. The BLM gene cluster or catalytic domains can be used individually or collectively to produce thiazolifne, thiazolifne, bithiazoline and bithiazolime-containing microbial metabolites. The BLM gene cluster may also be used to produce
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BOPCaR I; bovine parathyroid calcium receptor; hyperparathyroidism;
                                                                                                                                                                                                                                                                                                    New bleomycin gene cluster components useful for peptide and/or polyketide metabolites, especially bleomycin, production and for chemically modifying biological molecules -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14; DB 21;
Pred. No. 8.7e+02;
0; Mismatches 0;
                                                                                                                                                                                       Edwards DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18 BP; 5 A; 6 C; 5 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAQ88809 standard; cDNA to mRNA; 32 BP.
                                                                                                                                                                                    Chen M,
                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 22; 162pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  70.0%;
06-JAN-1999; 99US-0115435.
05-FEB-1999; 99US-0118848.
05-JAN-2000; 2000US-0477962.
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93WO-US01642.
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/mod_base=
                                                                                                                                                                                    Du L, Sanchez C,
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nes 14; Conservative
                                                                                                                      (REGC ) UNIV CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17
                                                                                                                                                                                                                                            WPI; 2000-465974/40.
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The invention provides an erythromycin-producing organism that expresses a heterologous oxygen-binding protein. The oxygen-binding protein can be used in a method for increasing erythromycin production that comprises culture of oxygen-binding protein under erythromycin production that comprises saccharopolyspora erythraea with an exogenous DNA construct, comprising directing integration of the construct at the phi C31 attachment site of the S. erythraea chromosome. The organism and the methods are used to provide more efficient production of erythromycin. Sequences AAX05608-609 represent PCR primers used for the amplification of PmerR promoter. This is used in the construction of expression vectors containing Vitreoscilla haemoglobin (Vhb) and chromosomal integration of the Vhb expression cassette in S. erythraea.
frames and other nucleic acid sequences from the genome of Chlamydia pneumoniae (see AAX91990). C. pneumoniae causes respiratory diseases such as pneumonia and bronchitis and is thought to be a contributing factor in heart disease, sarcoidosis, sinustis, purlent otitis media, erythema nodosum or pharyngitis. The polypeptides encoded by the open reading frames of the C. pneumoniae genome (see AAX3684-AX3679) can be used in immunogenic compositions as vaccines. Vectors containing C. pneumoniae nucleotides sequences can also be used as immunogenic compositions, especially where the vector directs the expression of a neutralising epitope of C. pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Erythromycin; oxygen-binding protein; transformation; S. erythraea;
PmerR promoter; Vitreoscilla; haemoglobin; Vhb; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New organism expressing oxygen-binding protein - useful, e.g. methods for highly efficient production of erythromycin
                                                                                                                                                                                                                                 Length 20;
                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                 67.0%; Score 13.4; DB 20;
93.3%; Pred. No. 1.7e+03;
ive 0; Mismatches 1;
                                                                                                                                                                                   Sequence 20 BP; 6 A; 4 C; 7 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 26 BP; 6 A; 9 C; 3 G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PmerR promoter amplifying reverse primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kallio P, Minas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 15; 44pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97US-0861450
                                                                                                                                                                                                                                                                                                                                                                                                                 AAX05609 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                 14; Conservative
                                                                                                                                                                                                                                                                                                  1 attectegetgetgg 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bailey JE, Brunker P,
                                                                                                                                                                                                                                                                                                                               19 ACTCCTCGCTGG 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (SOLI-) SOLIDAGO AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1999-070154/06
                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Streptomyces sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-MAY-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-NOV-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                AAX05609;
                                                                                                                                                                                                                                                                                                                                                                                  RESULT 10
AAX05609/c
                                                                                                                                                                                                                                                                Matches
   88888888888888
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Respiratory disease; pneumonia; bronchitis; heart disease; sarcoidosis; sinusitis; purulent otitis media; erythema nodosum; pharyngitis; vaccine; neutralising epitope; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAX91991-X97517 represent PCR primers used to amplify open reading
                                                                                                                                                                                                                                                              AAQ88809 was used in combination with AAQ88810 as primers for the PCR amplification of BoPcaR I, bovine parathyroid calcium receptor, which was used to test the effectivess of new calci-mimetics that mimics the action of extracellular ca ions. These calci-mimetics can be used in the treatment of a variety of diseases associated with abnormal levels of ca in cells, blood and plasma, specifically hyperparathyroidism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                 Compsn. contg. partly new calci-mimetic and calcilytic cpds. -
for treating parathyroidism, Paget's disease etc. and for
diagnosis, also new ion receptors and associated nucleic acid,
antibodies and transgenic animals
                                                                  Hebert SC;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR primer used to amplify an ORF of Chlamydia pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                  Fuller FH,
                                                                                                                                                                                                                                                                                                                                                                                                                                            68.0%; Score 13.6; DB 15; 70.0%; Pred. No. 1.4e+03; ive 2; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                Sequence 32 BP; 5 A; 4 C; 8 G; 8 T; 7 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Genome sequence of Chlamydia pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Page 1831; Disclosure; 1912pp; English
                                                                  Del MAR EG,
                                                                                                                                                                                                                                 Disclosure; Page 100; 283pp; English.
                  (BGHM ) BRIGHAM & WOMENS HOSPITAL (NPSP-) NPS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВР
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 attectegetgetggaa 20
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                                                                                Van WAGENEN BC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 70.03
Matches 14; Conservative
                                                                  Brown EM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAX96509 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chlamydia pneumoniae.
                                                                                                                WPI; 1994-293958/36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-357842/30
                                                                                                                                               contg.
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                                                                  Balandrin MF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-NOV-1998;
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                                                                                   Wemeth EF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAX96509;
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Gaps

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Indels

3;

Mismatches

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Matches 15; Conservative

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAX09121-X10268 are allele-specific oligonucleotide primers used in the isolation of various biallelic polymorphic markers found in the human genome (represented in AAX10269-X12937). These primers can be used in a method for determining polymorphic forms in an individual for use in e.g. forensics, paternity testing or for phenotypic typing for diseases such as agammaglobulinemia, diabetes insipidua. Fesch-Wyhan syndrome, muscular dystrophy, Wiskott-Aidrich syndrome, Fabry's disease, familial hypercholesterolemia, polycystic kidney disease, hereditary haemorrhagic telanglectasia, familial colonic polyposis, Ehlers-Danlos syndrome, osteogenesis imperfecta, acute intermittent porphyria,
                                                                                                                                                                                                                                                                                                                                      Polymorphism; biallelic; human; forensic; paternity testing; disease; detection; phenotypit typing; characteristic; infection; hereditary; autoimmune disease; cancer; inflammation; drug; therapy; medicament; treatment; marker; primer; ss.
                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated nucleic acid segments from the human genome - used for determining polymorphic forms for use in e.g. forensics, paternity testing or phenotypic typing for disease
                                        ;
0
   Length 26;
                                                                                                                                                                                                                                                                                                       Human biallelic polymorphic marker downstream primer #144.
                                      Indels
 Score 13.4; DB 20;
Pred. No. 1.7e+03;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 8 A; 4 C; 7 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (WHED ) WHITEHEAD INST BIOMEDICAL RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 16; Page 63; 310pp; English.
                                      ó;
                                                                                                                                                                                                  BP.
   67.0%;
93.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             prophylaxis of such diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    97WO-US20313.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       96US-0030455
                                                                                                                                                                                               AAX09838 standard; DNA; 20
                                                                                                                                                                                                                                                                      (first entry)
                                  14; Conservative
                                                                       6 tcgctgctggtggaa 20
                                                                                         26 TAGCTGCTGGTGGAA 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1998-286974/25.
Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-NOV-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                   AAX09838;
                                    Matches
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Score 13.2; DB 19; Length 20; Pred. No. 2.1e+03;

66.0%; 83.3%;

Query Match Best Local Similarity

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The invention relates to rice 5-enolpyruvylshikimate phosphate synthase (EPSPS) genomic DNA (AAC87188). The invention also relates to an expression cassette comprising, in the 5'-3' direction, one or more transcriptional enhancer elements selected from AAC87190-C87196), the rice EPSPS promoter, genomic DNA encoding a EPSPS chloroplast transit peptide, genomic DNA encoding a EPSPS chloroplast it is resistant to glyphosate (AAC87189), and a transcriptional terminator. The glyphosate resistant EPSPS contains a region (AAB29793) containing two amino acid substitutions relative to the corresponding wild type region (AAB29792). The invention also encompasses plant genomic EPSPS sequences identified via screening with a rice EPSPS intronic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequence; vectors and host plant cells comprising a nucleic acid sequence of the invention; transgenic plants (and tissues and seeds thereof) comprising a nucleic acid sequence of the invention, optionally further transformed with a DNA encoding an insect, fungal, viral, bacterial, nematode, stress or herbicide resistance protein; and methods of producing the transgenic plants of the invention. The nucleic acids and constructs of the invention are used to produce a wide variety of morphologically normal, glyphosate resistant plants. The glyphosate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            maize
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel polynucleotide encoding 5-enolpyruvylshikimate phosphate synthase, used to produce transgenic plants e.g. banana, wheat, or rice, having resistance or tolerance to glyphosate herbicide
                                                                                                                                                                                                                                                             Rice EPSPS; 5-enolpyruvylshikimate phosphate synthase; glyphosate resistance; herbicide resistance; transgenic plant; expression construct; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pickerill AP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bachoo S,
                                                                                                                                                                                                                               Rice mutant EPSPS PCR primer, SEQ ID NO:40.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 20; Page 51; 87pp; English.
                                                                                                                     AAC87185 standard; DNA; 23 BP.
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99GB-0030213.
99GB-0017839.
99GB-0017840.
99GB-0017846.
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99GB-0030207
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2 ttcctcgctgctggtgga 19
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                                20 TICCICICITCICGIGGA
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                                                                                                                                                                                                                                                                                                                                sativa.
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21-DEC-1999,
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                                                                                                                                                                                                                                                                                                                                                   Synthetic
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Gaps

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Indels

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Mismatches

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100.0%; Pred. No. 2.6e+03;

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13; Conservative
   Best Local Similarity
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29-JUL-1999;
21-DEC-1999;
21-DEC-1999;
21-DEC-1999;
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29-APR-1999;
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                                                                                                                                                                                                                                       Synthetic.
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              Matches
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resistant plants produced are particularly maize, soybean, cotton, sugarbeet and canola, but also other field crops, fruits and vegetables, turf and forage grasses and nut-producing plants. The plants are optionally resistant to insects, fungi, viruses, bacteria, nematodes, stress, desiccation and/or other herbicides. They can be used in the production of a herbicides for the high throughput in vitro screening of potential herbicides. The present sequence represents a PCR primer used in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel polynuclectide encoding the rice 5-enolpyruvylshikimate phosphate synthase, used to produce glyphosate tolerant or resistant plants -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to a glyphosate resistant rice 5-enolpyruvylshikimate phosphate synthase (EPSPS) gene. This gene can be used to produce plant tissue and/or morphologically normal fertile whole plants which are tolerant or resistant to glyphosate herbicide, and in the production of a herbicidal target for the high throughput in vitro screening of potential herbicides.
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0
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                                                                                                                                                                                                                                                                                                                                               Glyphosate; 5-enolpyruvylshikimate phosphate synthase; EPSPS; herbicide resistance; ss.
                                                                                                                                    Length 23;
                                                                                                                                                         Indels
                                                                                                                                  Query Match 65.0%; Score 13; DB 21; I Best Local Similarity 100.0%; Pred. No. 2.6e+03; Matches 13; Conservative 0; Mismatches 0;
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                                                                                                  Sequence 23 BP; 5 A; 7 C; 5 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 23 BP; 5 A; 7 C; 5 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Andrews CJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 20; Page 50; 85pp; English.
                                                                                                                                                                                                                                                          AAC88395 standard; DNA; 23 BP.
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99GB-0009972.
99GB-0017837.
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99GB-0030206.
99GB-0030214.
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                                                                                                                                                                               8 gctgctggtggaa 20
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21-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                               AAC88395;
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                                                                                                                                                                                                                                     RESULT 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Isolated polynucleotide encoding a 5-enolpyruvylshikimate phosphate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              synthase from rice is used for producing transgenic plants with enhanced resistance to glyphosate herbicide -
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                                                                                                                                                                                                                                                                                                                                                  5-enolpyruvylshikimate phosphate synthase; EPSPS; herbicide resistance; glyphosate; ss.
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100.0%; Pred. No. 2.6e+03;
iive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Andrews CJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 20; Page 49; 98pp; English
                                                                                                                                                                 BP.
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99GB-0017836.
99GB-0017843.
99GB-0030202.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99GB-0009969
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99GB-0030212
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Best Local Similarity 100.
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                          Primer universal reverse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hawkes TR, Warner SAJ,
                                                                                                                                                                 AAC89319 standard; DNA;
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                                           GCTGCTGGTGGAA 8
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||||||||||||||
    gctgctggtggaa
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RESULT 15 AAA82953/c

DB 21; Length 23;

65.0%; Score 13;

Query Match

AAA82953 standard; DNA; 19 BP

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The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinases other than cell-cycle dependent kinases CDK1. PCNA and Cyclin B1. Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restences by introduction of the ribozyme is resistant to endonuclease activity and hence is efficient in restenosis treatment.
                                                                                                                                                                                                                                                                                                                                                                                  New hairpin and hammerhead ribozyme for inhibiting restenosis, cleaves RNA encoding a cyclin or cell-cycle dependent kinase other than CDK1, PCNA and Cyclin B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 12.8; DB 21; Length 19;
Pred. No. 3.2e+03;
0; Mismatches 2; Indels (
                                                                                                                Ribozyme; hairpin; hammerhead; gene therapy; vasotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 3 A; 8 C; 5 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                 Robbins JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 54; 109pp; English.
                                                                                                                                                                                                                                                                                                                                 Barber JR,
                                                                                    cdk6 ribozyme binding site #13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 64.0%;
Best Local Similarity 87.5%;
Matches 14; Conservative
                                                                                                                                                                                                                                           99WO-US28772.
                                                                                                                                                                                                                                                                        98US-0110954
                                                       04-DEC-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                 Welch PJ,
                                                                                                                                                                                                                                                                                                                                                            WPI; 2000-412314/35.
                                                                                                                                                                                                                                                                                                    (IMMU-) IMMUSOL INC
                                                                                                                                                                                    WO200032765-A2.
                                                                                                                                 restenosis; ss
                                                                                                                                                                                                                                                                        04-DEC-1998;
                                                                                                                                                                                                                                             06-DEC-1999;
                                                                                                                                                                                                                08-JUN-2000.
                                                                                                                                                          Mammalia.
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                           AAA82953;
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Search completed: October 2, 2001, 16:18:45 Job time: 15489 sec

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Searched:

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Minimum Maximum Database

Result Š.

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GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REPERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310B
CURRENT APPLICATION NUMBER: US/09/377,310B
CURRENT PILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 16
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
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US-08-049-283A-22
US-08-881-037-81
US-07-796-106-16
US-08-418-859-31
US-08-643-181-31
US-08-643-181-32
US-08-943-529-1
US-08-943-529-1
US-08-943-529-1
US-08-050-132A-11
US-08-050-132A-11
US-08-050-132A-11
US-08-050-132A-11
US-08-050-132A-17
US-08-050-132A-17
US-08-050-132A-17
US-08-050-132A-17
US-08-050-132A-17
US-08-050-134-17
US-08-050-134-17
US-08-928-881-12
US-08-928-881-12
US-08-928-881-12
US-08-928-881-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CTHER INFORMATION: antisense sequence US-09-377-310-16
                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 16, Application US/09377310B
; Patent No. 6133031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.08; Fr. 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 100.
Best Local Similarity 100.
Matches 20; Conservative
    0000000000000000000
    RESULT 2
US-09-377-310-36
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Sequence 36, Appl
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9.071 Million cell updates/sec
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in or equal to the score of the result being printed,
analysis of the total score distribution.
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Sequence 27:
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    /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
    /cgn2_6/ptodata/1/ina/5B_COMB.seq:*
    /cgn2_6/ptodata/1/ina/6A_COMB.seq:*
    /cgn2_6/ptodata/1/ina/6B_COMB.seq:*
    /cgn2_6/ptodata/1/ina/PcTUS_COMB.seq:*
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Compugen Ltd.
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US-09-377-310-36
US-09-168-186-87
US-09-16-186-87
US-09-16-186-87
US-09-17-389-275
US-08-17-389-275
US-08-475-228A-275
US-08-475-228A-275
US-08-049-283A-21
US-08-17-389-285
US-08-475-228A-285
US-08-477-890-2
US-08-477-890-2
US-08-467-034A-2
US-08-467-034A-2
US-08-467-034A-2
US-08-467-034A-2
US-08-467-133-2
US-08-468-11-33-2
US-08-30-11-33-2
US-08-30-11-33-2
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US-07-960-982-23
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Copyright (c) 1993 - 2000
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Maximum Match 100%
Listing first 45 summaries
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                                                                         OM nucleic - nucleic search, using sw model
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US-09-166-186-87
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                                                                                                                                                                                            DB 3; Length 15; 46;
                                                                                     0; Indels
                                                          75.0%; Score 15; DB 100.0%; Pred. No. 46; tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRINCE APPLICATION NUMBER: US/08/585,684B
FILING DATE: January 16, 1996
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 60/000,951
FILING DATE: July 7, 1995
ATTORREY AGENT INFORMATION:
NAME: WALDURY, RICHARD
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/078
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECHONE: (213) 489-160
TELECHONE: (213) 955-0440
TELERAX: (213) 955-0440
TELERAX: 67-3510
SEQUENCE CHARACTERISTICS:
   ; OTHER INFORMATION: antisense sequence US-09-377-310-36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 ctaggggaggctcagtgt 18
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                                                           Query Match 75.0
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 18 base pairs
TYPE: nucleic acid
                                                                                                             3 aggggaggctcagtg 17
                                                                                                                                     1 aggggaggctcagtg 15
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Best Local Similarity
Matches 15; Conserv
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US-08-585-684B-2592/C
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US-08-585-684B-2592
FEATURE:
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RESULT

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Sequence 87, Application US/09166186A

Sequence 87, Application US/09166186A

GENERAL INFORMATION:
APPLICANT: BAKEr, Erenda
APPLICANT: Bennett, C. Frank
APPLICANT: Bennett, C. Frank
APPLICANT: Shanahan; William R.
TITLE REFERENCE: ISHH-0322
CURRENT PALLICATION NUMBER: US/09/166,186A
CURRENT PILING DATE: 1998-10-05
NUMBER OF SEQ ID NOS: 250
SEQ ID NO 87
LENGTH: 20
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Sequence 2592.3.4 Application US/09038073
Patent No. 6194150
GENERAL INFORMATION:
APPLICANT: Stinchcomb, Daniel T.
APPLICANT: Jarvis, Thale
APPLICANT: Maswiggen, James
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE
TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES
NUMBER OF SEQUENCES: 2751
CORRESPONDENCE: 2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 13.2; DB 4;
Pred. No. 3.4e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastESQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/038,073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/585,684
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/078
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELERAX: (213) ---
TELERAX: 67-3510
TELERA: 67-3510
; INFORMATION FOR SEQ ID NO: 2592:
SEQUENCE CHARACTERISTICS:
TOWNTH: 18 base pairs
TOWNTH: 1.1 acid
                                                                                                                                                                                                                                                    ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
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83.3%;
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Best Local Similarity 83.3
Matches 15; Conservative
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STATE: California
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Score 13.2; DB 1; Length 47;
Pred. No. 3.6e+02;
0; Mismatches 3; Indels
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US-08-171-389-275
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COMPUTER: IBM PC compatible
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURREMY APPLICATION DATA:
APPLICATION NUMBER: US/08/123,936
   PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Edwards, Cynthia A.
APPLICANT: Cantor, Charles R.
APPLICANT: Cantor, Charles R.
APPLICANT: Andrews, Beth M.
APPLICANT: Turin, Lisa M.
TITLE OF INVENTION: Screening Assay for the TITLE OF INVENTION: DNA Binding Molecules NUMBER OF SEQUENCES: 640
CORRESPONDENCE ANDRESS:
ADDRESSEE: Genelabs Technologies, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 33,875
HR: 4600-0175/G19P3
                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 07/996,783
FILING DATE: 23-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/723,618
FILING DATE: 27-JUN-1991
PRIOR APPLICATION NUMBER: US 08/081,070
FILING DATE: 22-JUN-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/123,936
FLIING DATE: 17-SEP-1993
PRIOR APPLICATION DATA:
                                                         US/08/171,389
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Fabian, Gary R.
REGIESTRATION NUMBER: 33,875
REFERENCE/DOCKET NUMBER: 46
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: DNA (genomic) HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (415) 324-0860
TELEFAX: (415) 324-0960
INFORMATION FOR SEQ ID NO: 275
SEQUENCE CHARACTERISTICS:
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83.3%;
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                                 CURRENT APPLICATION DATA APPLICATION NUMBER: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 47 base pairs
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EDNESS: double
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MEDIUM TYPE: Floppy
                                                                                                                     CLASSIFICATION: 435
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Best Local Similarity
Matches 15; Conserv
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USA
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COUNTRY: USI
ZIP: 94063
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                                                                                       FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
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CITY: Re
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                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 87, Application US/09313932A; Patent No. 6228642; GENERAL INFORMATION: APPLICANT: Barent APPLICANT: Barert, C. Frank; APPLICANT: Benett, C. Frank; APPLICANT: Benett, C. Frank; APPLICANT: Bancher, Madeline M. APPLICANT: Shanahan, William R.; TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE MODULATION OF TNF-1TLE OF INVENTION: EXPRESSION; TITLE OF INVENTION: EXPRESSION; CURRENT APPLICATION NUMBER: US/09/313,932A; CURRENT APPLICATION STORE OF THE O
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                                                                                                                                                                                                       Length 20;
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Pred. No. 3.5e+02;
); Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Edwards, Cynthia A.
APPLICANT: Cantor, Charles R.
APPLICANT: Andrews, Beth M.
APPLICANT: Turin, Lisa M.
APPLICANT: Fry, Kirk E.
TITLE OF INVENTION: Sequence-Directed DNA Binding
TITLE OF INVENTION: Molecules, Compositions and Methods
NUMBER OF SEQUENCES: 641
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                    Score 13.2; DB 3;
Pred. No. 3.5e+02;
0; Mismatches 3;
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                                                                                 CTHER INFORMATION: antisense sequence US-09-166-186-87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 275, Application US/08171389 Patent No. 5578444 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             505 Penobscot Drive
TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                    66.0%;
83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Synthetic US-09-313-932-87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 66.0%;
Best Local Similarity 83.3%;
Matches 15; Conservative
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                       3 aggggaggctcagtgtgg 20
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OPERATING SYSTEM:
                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 15; Conserv
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LENGTH: 20
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APPLICANT: Edwards, Cynthia A.
APPLICANT: Cantor, Charles R.
APPLICANT: Andrews, Beth M.
APPLICANT: Andrews, Beth M.
APPLICANT: Fry, Kirk E.
TITLE OF INVENTION: Sequence-Directed DNA Binding
TITLE OF INVENTION: Molecules, Compositions and Methods
NUMBER OF SEQUENCES: 664
CORRESPONDENCE ADDRESS:
                             APPLICATION NUMBER: US 08/081,070
FILING DATE: 22-UN-193
ATTORNEY/AGENT INFORMATION:
NAME: Stratford, Carol A.
REGISTRATION NUMBER: 34,444
REPRENENCE/DOCKET NUMBER: 4600-0175.21/G19P3D2
TELECOMMUNICATION INFORMATION:
TELEFAX: (415) 324-0860
INFORMATION FOR SEQ ID NO: 275:
SEQUENCE CHARACTERISTICS:
LENGTH: 47 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
FILING DATE: 07-JUN 1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Genelabs Technologies, Inc.
STREET: 505 Penobscot Drive
CIIY: Redwood City
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ...and DATE: 23-DEC 1992
PRIOR APPLICATION DATA:
PILING
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FILING DATE: 20-DEC-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                UMBER: US 08/123,936
17-SEP-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Sequence 275, Application US/08482080A; Patent No. 6010849
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83.3%; Pred
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HYPOTHETICAL: NO
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Best Local Similarity 83.3
Matches 15; Conservative
         PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                           nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-482-080A-275/C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: U. ZIP: 94063
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Edwards, Cynthia A. APPLICANT: Edwards, Cynthia A. APPLICANT: Cantor, Charles R. APPLICANT: Andrews, Beth M. APPLICANT: Turin, Lisa M. APPLICANT: Turin, Lisa M. APPLICANT: Turin, Lisa M. APPLICANT: Turin, Lisa M. APPLICANT: Fry, Kirk E. TITLE OF INVENTION: Sequence-Directed DNA Binding TITLE OF INVENTION: Molecules, Compositions and Methods NUMBER OF SEQUENCES: 664 CORRESPONDENCE ADDRESS: ADDRESSE: Genelabs Technologies, Inc. STREET: 505 Penobscot Drive CITY: Redwood City STREET: CA COUNTRY: USA ZTATE: CA COUNTRY: USA ZTATE: CA COUNTRY: USA ZIP: 94063 COMPUTER READABLE FORM: MEDIUM TRABBABLE FORM: MEDIUM TRABBABLE FORM: COMPUTER: IBM PC COMPATIBLE OF PRATICE COMPUTER: IBM PC COMPATIBLE COMPATI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          66.0%; Score 13.2; DB 1; Length 47; 83.3%; Pred. No. 3.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HYPOTHETICAL: NO

ORIGINAL SOURCE:

INDIVIDUAL ISOLATE: Human estrogen-responsive gene pS2
105-08-123-936-278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3; Indels
                                                                                                                                                                                                                                                    NAME: Fabian, Gary R.
REGISTRATION NUMBER: 33,875
REESERENCE/DOCKET NUMBER: 4600-0075.32/G19P2
RELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 324-0880
TELEPA: (415) 324-0860
INFORMATION FOR SEQ ID NO: 275:
SEQUENCE CHARACTERISTICS:
LENGTH: 47 base pairs
TENGTH: 47 base pairs
TENGTH: 47 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/475,228A FILING DATE: 06-JUN-1995
PRIOR APPLICATION DATE: 08-JUS-396
FILING DATE: 17-SEP-1993
PRIOR APPLICATION DATA: APPLICATION NUMBER: US 07/996,783
FILING DATE: 23-DEC-1992
FILING DATE: 23-DEC-1992
PRIOR APPLICATION DATA:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/996,783
FILING DATE: 23-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/723,618
FILING DATE: 27-UN-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 07/723,618 FILING DATE: 27-JUN-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 275, Application US/08475228A Patent No. 5869241 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDED TO THE TOPOLOGY: Innear MOLECULE TYPE: DNA (genomic)
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Matches 15; Conservative
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Patent No. 5502176
GENERAL INFORMATION:
APPLICANT: Tenen, Daniel G.
APPLICANT: Pahl, Heike L.
APPLICANT: Burn, Timothy C.
TITLE OF INVENTION: Cell Specific Promoter and Uses Thereof NUMBER OF SEQUENCES: 34
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 12.8; DB 1; Length 30; Pred. No. 5.5e+02; 0; Mismatches 2; Indels
                                                                                                                                                                                                                                    Length 47;
                                                                                                                      Human estrogen-responsive gene pS2
                                                                                                                                                                                                                                                                                             3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURENT APPLICATION DATA:
APPLICATION NUMBER: U8/08/049,283A
FILING DATE: 14-APR-1993
CLASSIFICATION AFF.
APPLICATION NUMBER: 08/020,465
FILING APPLICATION DATA:
APPLICATION NUMBER: 07/837,776
FILING DATE: 19-FEB-1993
CLASSIFICATION NUMBER: 10-FEB-1992
CLASSIFICATION NUMBER: 22,592
FRIOK APPLICATION NUMBER: B1491-03'A
TTORNEY/AGENT INFORMATION:
NAME: B1000k, David E.
REGISTRATION NUMBER: B1491-03'A
TELECOMMUNICATION INFORMATION:
T
                                                                                                                                                                                                                                 Score 13.2; DB 5;
Pred. No. 3.6e+02;
0; Mismatches 3;
                           MOLECULE TYPE: DNA (genomic) HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA (genomic)
                                                                                                                                                                                                                                    66.0%;
83.3%;
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Best Local Similarity 87.5%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                       2 taggggaggctcagtgtg 19
                                                                                                                                                                                                                                                                                                                                                                                         22 TAGGGCAGGCTCTGTTTG 5
                                                                                                                                                                                                                                                            Best Local Similarity 83.3
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4 ggggaggctcagtgtg 19
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Massachusetts
                                                                                     ORIGINAL SOURCE:
; INDIVIDUAL ISOLATE:
PCT-US93-12388-275
   TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE:
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US-08-049-283A-21/c
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STATE: MZ
COUNTRY:
                                                                                                                                                                                                                                       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               HYPOTHETICAL: NO ORIGINAL SOURCE: INDIVIDUAL ISOLATE: Human estrogen-responsive gene pS2 US-08-482-080A-275
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Sequence-Directed DNA Binding
TITLE OF INVENTION: Molecules, Compositions and Methods
NUMBER OF SEQUENCES: 641
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genelabs Technologies, Inc.
STREET: 505 Penobscot Drive
CITY: Redwood City
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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PRIOR APPLICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 08/123,936
FILING DATE: 17-SEP-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/996,783
FILING DATE: 23-DEC-1992
ATTORNEY FAGENT INFORMATION:
NAME: Fabian, Gary R.
REGISTRATION NUMBER: 4600-0175.41/G19PCT2
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECHOME: (415) 324-0880
TELECHOME: (415) 324-0860
INFORMATION FOR SEQ ID NO: 275:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SCENARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/12388
                                                                               NAME: Brady, John F.
REGISTRATION NUMBER: 39,118
REFERENCE/DOCKET NUMBER: 4600-0175.20/G19P3D1
TELECOMNUNICATION INFORMATION:
TELEPHONE: (650) 324-0880
TELEPHONE: (650) 324-0860
INFORMATION FOR SEQ ID NO: 275:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
APPLICATION NUMBER: US 08/081,070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCT-US93-12388-275/c; Sequence 275, Application PC/TUS9312388; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: DNA (genomic) HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               66.0%;
83.3%;
                           FILING DATE: 22-JUN-1993 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 taggggaggctcagtgtg 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 83.39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22 TAGGGCAGGCTCTGTTTG 5
                                                                                                                                                                                                                                                                                                                   LENGTH: 47 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 47 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                           linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
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                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY:
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Length 47;
                                                                                                                          APPLICANT: Edwards, Cynthia A.
APPLICANT: Cantor, Charles R.
APPLICANT: Andrews, Beth M.
APPLICANT: Turin, Lisa M.
TITLE OF INVENTION: Screening Assay for the Detection of WIMBER OF SEQUENCES: 640
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/123,936
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 12.8; DB 1;
Pred. No. 5.7e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4600-0075.32/G19P2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; INDIVIDUAL ISOLATE: Human SAA1 beta gene US-08-123-936-285
                                                                                                                                                                                                                                                                                                   Genelabs Technologies, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/996,783
FILING DATE: 23-DEC-1992
PRIOR APPLICATION NUMBER: US 07/723,618
FILING DATE: 27-UNH-1991
ATTORNEY/AGENT: 1FFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 285, Application US/08475228A
Patent No. 5869241
APPLICANT: Edwards, Cynthia A.
APPLICANT: Cantor, Charles R.
APPLICANT: Andrews, Beth M.
APPLICANT: Turin, Lisa M.
                                          US-08-123-936-285
; Sequence 285, Application US/08123936
: Patent No. 5726014
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                  STREET: 505 Penobscot Drive
CITY: Redwood City
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Fabian, Gary R.
REGISTRATION NUMBER: 33,875
REFERENCE/DOCKET NUMBER: 46
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 324-0880
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                64.0%;
87.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (415) 324-0960
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
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Best Local Similarity 87.5
Matches 14; Conservative
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STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM:
SOFTWARE: PatentI
                                                                                                                                                                                                                                                                                                                                                                                       USA
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US-08-475-228A-285
                                                                                                                                                                                                                                                                                                                                                                                                         94063
                                                                                                                                                                                                                                                                                                 ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                 STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH:
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                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Edwards, Cynthia A.
APPLICANT: Cantor, Charles R.
APPLICANT: Andrews, Beth M.
APPLICANT: Turin, Lisa M.
APPLICANT: Turin, Lisa M.
TITLE OF INVENTION: Sequence-Directed DNA Binding
TITLE OF INVENTION: Molecules, Compositions and Methods
NUMBER OF SEQUENCES: 641
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genelabs Technologies, Inc.
STREET: 505 Penobscot Drive
CITY: Redwood City
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 12.8; DB 1; Length 47;
Pred. No. 5.7e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/171,389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Fabian, Gary R. REGISTRATION NUMBER: 33,875
REFERENCE/DGCRET NUMBER: 4600-0175/G19P3
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; INDIVIDUAL ISOLATE: Human SAAl beta gene
US-08-171-389-285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 435
CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: US 08/123,936
FILING DATE: 17-SEP-1993
PRIOR APPLICATION NUMBER: US 07/996,783
PRIOR APPLICATION NUMBER: US 07/996,783
FILING DATE: 23-DEC-1992
PRIOR APPLICATION UDMAER: US 07/723,618
FILING DATE: 27-UN-1991
PRIOR APPLICATION UDMAER: US 08/081,070
APPLICATION UNMBER: US 08/081,070
RILING DATE: 22-UN-1991
FILING DATE: 22-UN-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                   Sequence 285, Application US/08171389
Patent No. 5578444
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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MOLECULE TYPE: DNA (genomic)
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (415) 324-0880
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (415) 324-0960
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
LENGTH: 47 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 64.0%;
87.5%;
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MEDIUM TYPE: Floppy disk
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18 GCGGAGGCTCAGTGAG 3
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: U
ZIP: 94063
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE:
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STATE:
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Gaps

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8 AGGGAAGGCTCAGTAT 23

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TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
HYPOTHETICAL: NO
ORIGINAL SOURCE:
INDIVIDUAL ISOLATE: Human SAA1 beta gene
                                                                                                                                                                                   LENGTH: 47 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 14; Conserv
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64.0%; Score 12.8; DB 2; Length 47; illarity 87.5%; Pred. No. 5.7e+02; Conservative 0; Mismatches 2; Indels 3 aggggaggctcagtgt 18 |||| ||||||||| | 8 AGGGAAGGCTCAGTAT 23 δ g

Search completed: October 2, 2001, 16:03:51 Job time: 14595 sec

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Potyvirus P1 prote Reporter gene cons Oligo "g" used in Hygromycin phospho Primer Cypozbir us Sequence of 5'LH p CDIIb gene promote

Scoring table:

Searched:

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Database

Perfect score:

Title:

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Run

Sequence:

Recombinant retrov Fibrinogen B-beta primer for bet P1 protein cod

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Human Human

PNIV3606

Human hyperpolaris PCR primer #1 to a SH2 binding protei

Primer R2 used to

for Fv for Fv

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PCR

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Primer

Human NK2R

Minisatellite MS32 Minisatellite MS32 MS32 minisatellite

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New antisense compounds for inhibiting focal adhesion kinase expression, especially useful for inhibiting retinal neovascularization, or for diagnosing and treating e.g. colon cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; focal adhesion kinase; FAK; signal transduction; cancer; embryonic development disorder; angiogenic disorder; wound healing; antisense; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human focal adhesion kinase antisense sequence #14.
                                                                                                                                                                                                                                                                                                                                                                                     ALIGNMENTS
                                                                                 AAV19540
AAA10258
AAX33879
AAT58070
                                                                                                                          AAZ23620
AAC66808
AAX06060
AAX59277
AAX58529
AAV39103
AAQ25680
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AAA58779
AAC60289
AAC66789
AAQ34833
                              AAT04232
AAQ69535
AAT63997
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AAT35053
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AAZ50600
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 99US-0377310
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  (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaarde WA;
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Homo sapiens.
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RESULT
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Primer for human c
Human estrogen-res
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Test sequence from
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                                                                     Search time 876.95 Seconds (without alignments)
14.320 Million cell updates/sec
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          GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                          Fotal number of hits satisfying chosen parameters:
                                                                                                                                                                                         730101 seqs, 313950809 residues
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                                                                      2001, 16:18:43
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT18223
AAT18225
AAX67092
AAA40846
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Gapop 10.0 , Gapext 1.0
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HBEGF; heparin-binding epidermal-like growth factor; saporin; cytotoxic; ribosome-inactivating protein; targeted delivery; tumbreast cancer; bladder cancer; restenosis; therapy; primer; PCR; polymerase chain reaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conjugate comprising HBEGF polypeptide and targeted agent - use treat HBEGF-mediated patho:physiological conditions, e.g. solid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 6; Page 94; 163pp; English.
                                                                                                                                                                                                                                                                                                                                                                             95US-0441979.
94US-0305771.
                                                                                                                                                                                                                                                                                                                                                    95WO-US12205.
                                                                                             AAT18223 standard; DNA; 34
                                                                                                                                                                          HBEGF DNA antisense primer
                                                                                                                                                24-JUN-1996 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tumour and skin disorders
 3 aggggaggctcagtg 17
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Best Local Similarity
Matches 14; Conserv
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                                                                                                                                                                                                                                                                        Synthetic.
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                       The present invention describes a number of phosphorothioate antisense sequences to the human focal adhesion kinase (FAK) protein. This protein is involved in integrin-mediated signal transduction, and is implicated in cancer, particularly colon, breast and oral tumours, embryonic development disorders, angiogenic disorders and wound healing. The antisense sequences, including the one shown here, can be used in the treatment of all of these.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes a number of phosphorothioate antisense sequences to the human focal adhesion kinase (FAK) protein. This protein is involved in integrin-mediated signal transduction, and is implicated in cancer, particularly colon, breast and oral tumours, embryonic development disorders, angiogenic disorders and wound healing. The antisense sequences, including the one shown here, can be used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense compounds for inhibiting focal adhesion kinase expression, especially useful for inhibiting retinal neovascularization, or for diagnosing and treating e.g. colon cancer -
                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                               Human; focal adhesion kinase; FAK; signal transduction; cancer; embryonic development disorder; angiogenic disorder; wound healing; antisense; phosphorothloate; ss.
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                                                                                                                                                                          22; Length 20;
                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                      Human focal adhesion kinase antisense sequence #34.
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                                                                                                                                                                            DB 22
0.53;
                                                                                                                                  Sequence 20 BP; 3 A; 3 C; 10 G; 4 T; 0 other;
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                                                                                                                                                                                                     Mismatches
                                                                                                                                                                          Score 20;
Pred. No. (
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Claim 3; Column 23; 30pp; English.
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                                                                                                                                                                          100.0%;
100.0%;
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                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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treatment of all of these.
                                                                                                                                                                        Query Match
Best Local Similarity 100.
Matches 20; Conservative
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McDonald JR, Nova MP, Sosnowski BA;

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Sense and antisense PCR primers (AAT18222-23) were used to amplify DNA (see AAT17958) coding for human heparin-binding epidermal-like growth factor (HBEGF) (AAR92897) using p232B1 as template. The PCR product has a 5' NCOI site and a 3' stop codon followed by an SalI site. It was ligated into PETSAP-MCS (see AAT18220-21). The resulting plasmid (p236B1) (see AAT18232) codes for a fusion (AAR92921) of HBEGF and saporin linked by an Ala-Met-Ala linker. This was expressed in Escherichia coli and is useful for targeting cytotoxic saporin e.g. to solid tumour cells bearing the HBEGF
                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                   Length 34;
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Pred. No. 9e+02;
); Mismatches 1;
                                                                                                                                                                                                                                                                                                    Sequence 34 BP; 9 A; 8 C; 9 G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                67.0%;
93.3%;
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Gaps

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Indels

Length 15;

75.0%; Score 15; DB 22; I 100.0%; Pred. No. 1.4e+02; Live 0; Mismatches 0;

Conservative

Best Local Similarity Matches 15; Conserve

Query Match

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95WO-US15516
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
nes 15; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                             present invention.
                                                                                                                                                                               Beigelman L,
           22-NOV-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-AUG-2000
                                                                              20-APR-1995
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04-MAY-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                           HBEGF; heparin-binding epidermal-like growth factor; saporin; cytotoxic; ribosome-inactivating protein; targeted delivery; tumour; breast cancer; bladder cancer; restenosis; therapy; primer; PCR; polymerase chain reaction; baculovirus; ss.
                                                                                                                                                                                                                                                   - used to
                                                                                                                                                                                                                                                                                                          An antisense PCR primer (AAT18225) was used with sense primers AAT1824 or AAT18227) to amplify DNA (see AAT17958) coding for human heparin-binding epidermal-like growth factor (HBEGF) (AAR92897) using p232B1 as template. The PCR products were lighted binto pBLMBBacll1 (Invitrogen) (see AAT18228 and used for baculovirus expression of HBEGF. Conjugates of HBEGF with saporin are useful for targeted treatment of e.g.
                                                                                                                                                                                                                                                  Conjugate comprising HBEGF polypeptide and targeted agent - us treat HBEGF-mediated patho:physiological conditions, e.g. solid
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                                                                                                                                                                                                                                                                                                                                                                                                                                Length 35;
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Pred. No. 9e+02;
0; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 35 BP; 10 A; 8 C; 8 G; 9 T; 0 other;
                                                                                                                                                                                                                                                                                          Example 8; Page 96; 163pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAX67092 standard; RNA; 18 BP.
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93.3%;
                                                                                                                                       95WO-US12205.
                                                                                                                                                          95US-0441979.
         HBEGF DNA antisense primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                            Houston LL,
                                                                                                                                                                                                                                                                      tumour and skin disorders
                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 93.3
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 diagnosis; ss
                                                                                                                                                                                                                                                                                                                                                                               solid tumours
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9618736-A2
                                                                                               WO9608274-A2
                                                                                                                                       13-SEP-1995;
                                                                                                                                                           16-MAY-1995;
                                                                                                                                                                   13-SEP-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-JUL-1999
                                                                                                                   21-MAR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-JUN-1996
                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAX67092
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The present invention describes a novel enzymatic nucleic acid (ENA)

having a hammerhead motif (EM) comprising: (i) at least 5 ribose

residues; (ii) a 2'-C-allyl modification at position 4 of the ENA; (iii)

at least ten 2'-O-methyl modifications; and (iv) a 3'-end modification.

The ENA's can inhibit collagenase and stromelysin production in the particularly osteoarthritis or the treatment or prevention of arthritis, particularly osteoarthritis or rheumacoid arthritis. The ENA's can also be used to treat antigen presenting cells of adonor to induce tolerance in a recipient to an alloantigen of a donor. They can also be used for enhancing graft tolerance or for treating autoimmune disease, and for creating allergies and other inflammatory conditions. The ENA's can also be used in diagnosis. Ribozyme therapy impacts on the expression of stromelysin without introducing the non-specific effects upon gene expression which accompany treatment with retinoids and dexamethasone.

The concentration of ribozyme required to affect a therapeutic treatment is lower than that required of antisense molecules, and is highly expectific. The present sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         - nsed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Enzymatic nucleic acid molecules having a hammer-head motif - us for the treatment of arthritis, induction of graft tolerance or treatment of auto-immune diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 iggen J, Pavco P, Stinchcomb
odak A, Usman N, Burgin A;
Thompson JD, Wincott F;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human TNFalpha antisense oligonucleotide ISIS# 21706.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 BP; 3 A; 11 C; 2 G; 2 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    McSwiggen J,
A, Modak A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 10; Page 216; 307pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matulic-Adamic J, Jarvis T,
                                                                                                                                                                              95US-0426124.
95US-0432874.
95US-0434509.
95US-0000951.
95US-0541365.
94US-0354920.
                                                                                                          94US-0363254.
                                                                        94US-0363253
                                                                                                                                                                                                                                                                                                                                                                    95US-0512861
                                                                                                                                                                                                                                                                                                                              95US-0000974
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gustofson J, M
L, Karpeisky A,
                                                                                                                                                                                                                                                                                                                                                                                                                                            (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 ctaggggaggctcagtgt 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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99US-0322134

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The present sequence is a PCR primer for human clone AAP12616HY coding sequence (see AAC90427). The AAP12616HY coding sequence is useful for modulating the immune and inflammatory responses, and hematopoiesis; and in gene therapy for treating allergies (e.g. asthma), autoimmune disorders (e.g. arthritis or transplantation rejection), in tissue treatment of surgical incisions and ulcers, such as stomach or diabetic ulcers). The AAP12616HY coding sequence is thought to encode a chemokine
                                                                                                                                                                                                                                                                                                    New chemokine polypeptide from fetal liver-spleen, useful in gene therapy, especially for modulating hematopoiesis, or immune or inflammatory responses, e.g. for repairing tissues or treating autokimmune disorders
                                                                                                                                                                                                                                                                                                                                                                                        Example 3; Page 78; 92pp; English.
                                                                                                                                    25-MAY-2000; 2000WO-US14572
   chromosome 2q11; ss.
                                                                                                                                                                                                                                                                    WPI; 2001-041140/05.
                                                                                                                                                                                                                                      Ford J, Yeung G;
                                                                 WO200073320-A1
                                                                                                                                                                                                     (HYSE-) HYSEQ
                                    Homo sapiens.
                                                                                                                                                                   27-MAY-1999;
                                                                                                    07-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAT91995/C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This sequence represents an antisense oligonucleotide sequence which targets a region of the human tumour necrosis factor alpha (TNFalpha) nucleotide sequence. TNFalpha is an important cytokine that plays a role in host defence. It is produced mainly in macrophages and monocytes in response to infection, invasion, injury or inflammation. Overexpression of TNFalpha can result in disease states, particularly in infectious, inflammatory and autoinmune diseases. The invention relates to antisense oligonucleotides, such as that represented by the present sequence which are capable of modulating the TNFalpha gene expression. The oligonucleotides optionally have a phosphorothoate backbone, and may also optionally contain at least one 2'-0-methoxyethyl modification. The oligonucleotides are useful for modulating the expression of human coligonucleotides are useful for modulating the archeology response, reducing the blood glucose level in a human and treating a human having a disease or condition associated with TNFalpha in Casamples of diseases associated with TNFalpha include diabetes, inflammatory bowel diseases multiple sclerosis, pancreatitis, rheumatoid arthritis, infectious disease, multiple sclerosis, pancreatitis or allogaraft rejection.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; clone pl2616HY; immune response modulation; autoimmune disorder;
inflammatory response; hematopoiesis; gene therapy; allergy; PCR primer;
autoimmune disorder; tissue repair; wound healing; chemokine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                    Oligonucleotide for treating diseases associated with human tumour necrosis factor-alpha (TNFalpha) such as, diabetes and rheumatoid arthritis, comprises nucleotide sequence complementary to intron of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            function of a selected nucleic acid sequence in adipose tissue.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             66.0%; Score 13.2; DB 21; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
pancreatitis; atopic dermatitis; allograft rejection; autoimmune disease; inflammatory disease; ss.
                                                                                                                                                                                                                                                                    Shanahan WJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No. 1.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 5 A; 3 C; 9 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                     Example 6; Page 58; 283pp; English.
                                                                                                                                                                                                                                                                    Bennett CF, Butler MM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human clone p12616HY PCR primer #1.
                                                                                                                                                                                                                                                                                                                                                                                       nucleic acid encoding TNFalpha -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAC90428 standard; DNA; 21 BP
                                                                                                                                                                                 98US-0166186.
99US-0313932.
                                                                                                                                                   99WO-US23205
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nes 15; Conservative
                                                                                                                                                                                                                                   (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                      WPI; 2000-303808/26.
                                                                                 WO200020645-A1
                                                                                                                                                                                                     18-MAY-1999;
                                                                                                                                                  05-OCT-1999;
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                                                                                                                  13-APR-2000
                                                                                                                                                                                                                                                                    Baker BF,
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Matches
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Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; chemokine alpha-3; CK alpha-3; treatment; tumour;
bronic infection; leukaemia; T-cell mediated autoimmune disease;
parasitic infection; psortasis; asthma; allergy; haematopoiesis;
growth factor; angiogenesis; wound healing; PCR primer; ss.
                                                                                                                                          ö
                                                                                                Length 21;
                                                                                                                                            Indels
                                                                                                Score 13.2; DB 22;
Pred. No. 1.1e+03;
                                     Sequence 21 BP; 7 A; 7 C; 4 G; 3 T; 0 other;
                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Primer for human chemokine alpha-3 cDNA.
and is located on chromosome 2q11.
                                                                                                                                                                                                                                                                                                                                BP.
                                                                                                66.0%;
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                                                                                                                                                                                                                                                                                                                              AAT91995 standard; DNA; 33
                                                                                                                                                                                    3 aggggaggctcagtgtgg 20
                                                                                                                                                                                                                                                                                                                                                                                                                  08-APR-1998 (first entry)
                                                                                                                      Local Similarity 83.3
hes 15; Conservative
                                                                                                                                                                                                          19 AGTGCAGTCTCAGTGTGG
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                                                                                                     Query Match
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(GENE-) GENELABS TECHNOLOGIES INC.
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91US-0723618.
92US-0996783.
                                                                                                                                                                                                                66.0%;
83.3%;
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                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                        15; Conservative
                                                                                                                                                                                                                                                                             22 TAGGGCAGGCTCTGTTTG
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                                                                                                                                                                                                                             Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-JUN-1991;
                                                                                                                                                                                                                                                                                                                                                                                               17-MAR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27-JUN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-SEP-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US5578444-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-NOV-1996
                                                                                                                                                      AAQ69891).
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                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                     AAT63987/c
                                                                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                               δλ
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                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A DNA protein-binding assay is provided, useful for screening libraries of synthetic or biological cpds. for their ability to bind DNA test sequences. The assay is versatile in that any number of test sequences can be tested by placing the test sequence adjacent to a defined protein-binding screening sequence. Binding of mols. to these test sequences changes the binding characteristics
                                                                                                                                       The present sequence is a primer for the cDNA encoding human chemokine alpha-3 (CK alpha-3). CK alpha-3 can be used to treat tumours. chronic infections, leukaemia, T-cell mediated autoimmune diseases, parasitic infections, psoriasis, asthma and allergy. It can also be used to regulate hematopoiesis, stimulate growth factor activity, inhibit angiogenesis and promote wound healing.
                                                                        treat
                                                                                                                                                                                                                                                                                      ;
                                                                   Polynucleotide(s) encoding human chemokine alpha-3 - used to tumours, chronic infections, auto:immune diseases, parasitic infections, psoriasis, asthma etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA protein-binding assay; test sequence; screening sequence; promoter; target; TATA box; Herpes Simplex Virus; HSV; origin of replication; UL9; transcription factor; TFIID: ds.
                                                                                                                                                                                                                                                              Score 13.2; DB 18; Length 33; Pred. No. 1.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Turin LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence-directed DNA-binding molecules - useful in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human estrogen-responsive gene pS2 target region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fry KE,
                                                                                                                                                                                                                          Sequence 33 BP; 8 A; 14 C; 7 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pharmaceuticals and as molecular reagents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Edwards CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 28; Page 349; 587pp; English.
                                                                                                                   Example 3; Page 54; 69pp; English.
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                                                                                                                                                                                                                                                                                     ;
                                                                                                                                                                                                                                                                                                                                                                                              AAQ69525 standard; DNA; 47 BP.
                                                                                                                                                                                                                                                             66.0%;
(HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                93WO-US12388.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     92US-0996783
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                                                                                                                                                                                                                                                                                                           3 aggggaggctcagtgtgg 20
                                                                                                                                                                                                                                                                                                                          28 ACGGGAGGCTCATGGTG 11
                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Andrews BM, Cantor CR,
                       Su JY;
                                             WPI; 1997-480232/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1994-234711/28.
                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 15; Conserv
                       Ni J,
                                                                                                                                                                                                                                                                                                                                                                                                                                             01-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-SEP-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
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                      Li H,
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of the protein mol. to its cognate binding sequence. When such a mol. binds the test sequence, the equilibrium of the DNA:protein complexes is disturbed, generating changes in the concentration of free DNA probe. One application of this method is to eucaryotic general transcription factors (e.g. TFIID), where the target region is typically selected from DNA sequences adjacent to the binding site for the eucaryotic transcription factor. Numerous exemplary test sequences are given: the sequences in AAQ69251-731 and AAQ698800 correspond to promoter targets (typically, TATA box-contg. sites) for human genes and the sequences in AAQ69732-849 correspond to promoter targets for viral genes. The test sequences may also be randomly generated. DNA:protein interaction may be used for screening purposes, e.g. the Herpes Simplex Virus (HSV) origin of replication and UL9 (see AAQ69851-52, AAQ69855 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Duplex DNA; target region; binding characteristic; DNA binding protein; TFIID; transcription factor; binding site; inhibition; enhance; inherited genetic disorder; cancer; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The sequences given in AAT63713-4312 represent duplex DNA's which act as target regions in the method of the invention. The method for altering the binding characteristics of a DNA-binding protein to duplex DNA comprises contacting the duplex DNA with a small molecule which binds sequence-specifically to a target region, where, when the small molecule is bound to the target region, it is adjacent to, but not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Altering binding characteristics of DNA binding proteins to duplex DNA - by attaching specific small cpd. to target region close to the protein's binding site, useful in treatment of viral disease, cancer
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 47;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fry KE, Turin LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 13.2; DB 15;
Pred. No. 1.2e+03;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 47 BP; 12 A; 15 C; 13 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
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overlapping by more than 4 bp, a binding site for a DNA-binding protein. The small molecule is added at a concentration effective to alter the binding of the DNA binding protein, pref. TFIID, to its binding site on the duplex DNA. The binding of the small molecule may inhibit or compounds isolated using this method are potentially useful as therapeutic agents for treatment of any disease which involves a specific DNA sequence, e.g. cancer, or inherited genetic disorders etc. The method is suitable for screening large biological or chemical ibraries and allows determination of sequence-specific and relative affinities of known DNA-binding agents for different DNA sequences. The design of these duplex DNA's allows a single DNA; protein interaction to be used for screening sequence specific, or preferential, DNA binding proteins that recognise almost any possible sequence (see also AAT49539-
                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                a DNA-binding molecule
                                                                                                                                                                                                                                                                                                                                                                          ;
0
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                                                                                                                                                                                                                                                                                                                                         Length 47;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Determination of DNA sequence preference of a DNA-binding mo based on inhibition of binding of protein to oligonucleotide sequence attached to test sequence
                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Turin LM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Test sequence from human estrogenresponsive gene pS2.
                                                                                                                                                                                                                                                                                                                                   Score 13.2; DB 18;
Pred. No. 1.2e+03;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                  Sequence 47 BP; 12 A; 15 C; 13 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fry KE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 3; Columns 241-242; 270pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP.
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91US-0723618.
92US-0996783.
93US-0123936.
95US-0475228.
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83.3%;
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ID AAX17275 standard; DNA; 47
                                                                                                                                                                                                                                                                                                                                                                                                                                22 TAGGGCAGGCTCTGTTTG 5
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Best Local Similarity 83.3
Matches 15; Conservative
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17-SEP-1993;
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where the screening sequence binds to the DNA-binding protein with a binding affinity that is independent of the DNA sequence of the test sequence, and where the mixture of duplex DNA test oligonocleotidess concludes several test sequences; (ii) incubating the test molecule, the mixture of duplex DNA test oligonocleotides and the DNA-binding protein for a time sufficient to permit binding of the test molecule to test sequences in the duplex DNA; (iii) separating unbound test oligonocleotides from test oligonocleotides bound to binding protein; (iv) maplifying the unbound test oligonocleotides; (v) repeating steps (ii) to (iv); (vi) isolating the amplified test oligonocleotides; and (vii) sequencing the isolated test oligonocleotides. Test sequences ANAI7001-X17481 and AAX17600 correspond to promoter targets for human genes and test sequences AAX17482-X17599 correspond to promoter targets
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; variant thrombospondin 1; variant thrombospondin 4; SNP;
polymorphism; vascular disease; coronary artery disease; forensics;
myocardial infarction; atherosclerosis; stroke; venous thromboembolism;
pulmonary embolism; paternity test; ds.
                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /standard_name= "single nucleotide polymorphism"
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                                                                                                                                                                                                                                                                                                             Length 47;
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                                                                                                                                                                                                                                                                                                         Score 13.2; DB 20;
Pred. No. 1.2e+03;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human gene single nucleotide polymorphism #1377.
                                                                                                                                                                                                                                                           Sequence 47 BP; 12 A; 15 C; 13 G; 7 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                            0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAF96616 standard; DNA; 21 BP.
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                                                                                                                                                                                                                                                                                                         66.0%;
83.3%;
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16-AUG-2000; 2000US-0225724.
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                                                                                                                                                                                                                                                                                                                                                                           2 taggggaggctcagtgtg 19
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Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                              22 TAGGGCAGGCTCTGTTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-226749/23.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAF96616;
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                                                                                                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; hyperpolarisation-activated channel; HAC3; neuroleptic; nootropic; cerebroprotective; antimigraine; antiarrhythmic; gene therapy; pacemaker dysfunction; familial sinus rhythm disease; sick sinus syndrome associated with atrial fibrillation; sinus tachycardia; bradycardia; ventricular arrhythmia; bipolar disease; schizophrenia; central nervous system disorder; migraine; seizure;
present invention provides a method of diagnosing a vascular disease
                   in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myocardial infarction, coronary heart disease, stroke, peripheral vascular diseases, venous thromboembolism and pulmonary embolism. Single nucleotide polymorphisms (SNRs) are also useful in forensics, paternity testing, genetic analysis and phenotype correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel human hyperpolarization activated channel 3 polypeptide useful to identify hyperpolarization-activated cation channels modulators for treating familial sinus rhythm diseases, and ventricular arrhythmias -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to the coding sequence and polypeptide sequence for human hyperpolarisation activated channel 3 (HAC3) (see AAC66779 and AAB28175). HAC3 comprises an alpha-subunit of a cation channel, which forms upon hyperpolarisation, a cation channel with an additional HAC. Modulators of HAC activity are useful for treating various pacemaker dysfunctions such as familial sinus rhythm diseases, sick sinus syndrome associated with atrial fibrillation, sinus tachycardias and bradycardias as well as ventricular arrhythmias. The modulators are also useful for treating other disorders involving
                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human hyperpolarisation-activated channel HAC3 PCR primer #11.
                                                                                                                                                                                                                                                                                                                     Score 12.8; DB 22; Length 21;
Pred. No. 1.7e+03;
); Mismatches 2; Indels (
                                                                                                                                                                                                                                                       Sequence 21 BP; 5 A; 2 C; 12 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAC66790 standard; cDNA; 25 BP.
                                                                                                                                                                                                                                                                                                                       64.0%;
87.5%;
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                                                                                                                                                                                                                                                                                                                     Query Match 64.0
Best Local Similarity 87.5
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                        5 gggaggctcagtgtgg 20
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abnormal ion flux, e.g., memory and learning disorders, sleeping disorders, bipolar disease, schizophrenia, central nervous system (CNS) disorders such as migraines, hearing and vision problems, seizures, and as neuroprotective agents (e.g. to prevent stroke). The Human HAC3 coding sequence is useful for treating the disorders by gene thorapy. The present sequence is a PCR primer which can be used to isolate the human HAC3 coding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                An electrophoretic mobility shift assay (EMSA) was used to investigate DNA/protein interactions in the CD11b promoter. Nuclear extracts from U937 (myeloid) or HeLa (non-myeloid) cells were incubated with a 130 bp probe which extends from bp -90 to +40 of the CD11b promoter. In addition to the 130 bp probe, 30 bp double stranded oligonucleotides were used as probes (AAT14576-79). The sequence between bp -64 and -59 constitutes a consensus binding site for the transcription factor Sp1, so to investigate whether complexes formed in the EMSA represent Sp1 binding, a 97 bp DNA fragment of the tissue plasminogen promoter contg. 2 characterised Sp1 binding sites was used as a competitor. The complexes, but not the PU.1 complex, were competed by the addition of DNA contg. Sp1 binding sites, indicating that they represent Sp1 binding the CD11b promoter.
                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CD11b gene promoter directing gene expression pref. in myeloid cell: - directs expression in transient transfection assays in transgenic animals, used to identify factors regulating myeloid cell-specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     promoter; myeloid-specific; CD11b; Sp1; PU.1; transcription factor; binding; transcription start site; ss.
                                                                                                                                                                                                                                           ó
                                                                                                                                                                                                     Length 25;
                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CDIIb gene promoter nucleotides -78 to -49 (wt-78).
                                                                                                                                                                                                     Score 12.8; DB 21;
Pred. No. 1.7e+03;
0; Mismatches 2;
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                                                                                                                                                Sequence 25 BP; 4 A; 15 C; 2 G; 4 T; 0 other;
                                                                                                                                                                                                                                           0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                       AAT14576 standard; DNA; 30 BP
                                                                                                                                                                                                       64.0%;
87.5%;
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92US-0837776.
93US-0020465.
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                                                                                                                                                                                                                                                                               5 gggaggetcagtgtgg 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1996-179337/18
                                                                                                                                                                                                     Query Match
Best Local Similarity
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13-FEB-1992;
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This sequence, and the sequence represented by AAT04233 are amplification primers for pNIV3606 (which contains human histamine H1 receptor CDNA up to the fourth transmembrane region). The sequences shown in AAT04237, and AAR79965 represent human histamine H1 receptor CDNA and encoded receptor protein, and were constructed using AAT04234-6. Histamine H1 receptor is part of a family of histamine receptor proteins. Of these sequences the greatest homology was found with bovine histamine H1 receptor (overall identity of 82%, with the transmembrane regions alone having an identity of 86%). These receptors are coupled to different second messenger pathways via quanine nucleotide regulatory proteins (G proteins). This receptor transduces the signal through calculation ion mobilisation via an increase in intracellular inositol 1.4.5-triphosphate levels. Plasmids containing human histamine H1 receptor CDNA (pref. pNIV3604B) are used to transform mammalian cells (pref. CHO cells). These transformed cells express the receptor on the cell surface. The transformed cells can then be used to identify ligands that bind to the encoded protein, esp. for drug screening. This sequence is also used to detect complementary mRNA in the cell by hybridisation.
                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             a \alpha rorresponds to bases 1 to 48 of human histamine H1 DNA."
                                                                                                                                                                                                                                                                                                                                            Polymerase chain reaction; PCR; primer; amplify; histamine HI receptor; human; bovine; transformation; drug screening; pNIV3604B; calcium ion mobilisation; incositol 1,4,5-triphosphate; CHO cell; guanine nucleotide regulatory protein; G protein; hybridisation; ss.
                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New nucleic acid encoding human histamine H1 receptor - useful diagnostically and for screening receptor binding drugs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Varsalona F;
Length 30;
                                   Indels
Score 12.8; DB 17;
Pred. No. 1.8e+03;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Noyer M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 43 BP; 10 A; 17 C; 4 G; 12 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Moguilevsky N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 3; Page 39; 49pp; English.
                                                                                                                                                                                                                                                                                                          pNIV3606 amplification primer #1.
                                                                                                                                                                                                 AAT04232 standard; DNA; 43 BP.
Query Match
Best Local Similarity 87.5%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        93GB-0022353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        93GB-0022353
                                                                                                                                                                                                                                                                      10-JAN-1996 (first entry)
                                                                       4 ggggaggctcagtgtg 19
                                                                                            14..43
/*tag=
/note= "
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1995-157301/21.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Key
misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GB2283239-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
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  Length 43;
                          Indels
 Score 12.8; DB 16;
Pred. No. 1.8e+03;
); Mismatches 2;
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  64.0%;
87.5%;
                          14; Conservative
                                                2 tagggggaggctcagtg 17
                                                                         27 TTGGGGAGGCTCATTG 12
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Job time: 15488 sec
Query Match
Best Local Similarity
                          Matches
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Appl Appl Appl

Sequence Sequence

Sequence

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Sequence: Perfect

Run on:

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Sequence Sequence

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APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REPERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310B
CURRENT APPLICATION NUMBER: US/09/377,310B
SOFTWARE: PAPLICATION DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
ERQ ID NO 15
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310B
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: PatentIn Ver. 2.0
        PCT-US95-05423-7

US-09-268-1140-29

US-09-556-318-10

US-08-468-846-7

US-08-468-846-7

US-08-846-062-10

US-07-937-609-12

US-08-936-632B-38

US-08-936-632B-38

US-08-131-11-12

US-08-581-543-30

US-08-111-382-10

US-08-131-538-6

US-08-131-538-6

US-08-513-974B-150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 20; DB 3;
Pred. No. 0.035;
                                                                                                                                                                           US-08-450-384-8
US-08-450-384-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                        ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ), OTHER INFORMATION: antisense sequence US-09-377-310-15
                                                                                                                                                                                                                                                       RESULT 1
18-09-377-310-15
Sequence 15, Application US/09377310B
Patent No. 6133031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             78-09-377-310-35
; Sequence 35, Application US/09377310B
; Patent No. 6133031
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LEWGTH: 15
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 tgttggtttccaatcggacc 20
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Best Local Similarity
Matches 20; Conserv
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Query Match
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                                                                                    (without alignments)
9.071 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 129,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15,
                                                                         ; Search time 417.38 Seconds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Description
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Sequence
Sequence
Sequence
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                                                                                                                                                                                                                                                                                                                                   /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
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/cgn2_6/ptodata/1/ina/PcTuS_COMB.seq:*
/cgn2_6/ptodata/1/ina/PcTuS_COMB.seq:*
         GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-377-310-15
US-09-377-310-15
US-09-377-310-35
US-08-295-676A-8
US-08-295-676A-8
US-08-612-895A-29
US-09-093-293-29
US-09-093-293-29
US-09-1653-44-8
US-08-912-100-42
US-08-912-38-12
US-08-913-914-919
US-08-73-479-54
US-08-73-479-54
US-08-73-479-74
                                                                                                                                                                                                                   Fotal number of hits satisfying chosen parameters:
                                                                                                                                                                                                324599 segs, 94655562 residues
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                                                                         2001, 16:03:50
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Maximum Match 100%
Listing first 45 summaries
                                                     nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
                                                                                                                                                                                                                                                                                                                          Issued_Patents_NA:*
                                                                                                                                                                                                                                                                                                                                                                                                                            score greater than or equal to
and is derived by analysis of
                                                                                                               US-09-757-100B-15
20
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                                                                          October
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11.8
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11.1.8
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Length 20 Indels

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/label- oligonucleotide
/note- "synthetic 5' primer used to create
recognition sites"
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TITLE OF INVENTION: A method for production of proteins in TITLE OF INVENTION: yeast
CORRESPONDENCE: 21
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  67.0%; Score 13.4; DB 3; Length 30; 93.3%; Pred, No. 1.2e+02; Live 0; Mismatches 1; Indels
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ZIP: 220-0747

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC COMPATIBLE

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PATENTIN Release #1.0, Version #1.25 (EPO)

CURRENT APPLICATION DATES: US/08/295,676A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc_feature
LOCATION: 5..10
OTHER INFORMATION: /function= "KpnI restriction site"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Birch, Stewart, Kolasch and Birch STREEF: P.O. Box 747
TTTY: Falls Church STATE: VA
                PRIOR DATE: 11 AUGUSTAS)

PRIOR DATE: 11 AUGUSTAS)

PREING DATE: 11 AUGUSTAS)

PREGISTRATION NUMBER: 34,235

REGISTRATION NUMBER: 34,235

REFERENCE/DOCKET NUMBER: 45753

TELECOMMUNICATION INFORMATION:

TELEPHONE: 617-523-34(0)

TELEPHONE: 617-523-6440

INFORMATION FOR SEQ ID NO: 129: SEQUENCE CHARACTERISTICS: LENGTH: 30 base pairs

TYPE: nucleic acid

STRANDEDNESS: Gouble
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 4
US-08-295-676A-8/C
Sequence 8, Application US/08295676A
Patent No. 5677172
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 09-SEPT-1 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 30 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 tgttggtttccaatc 15
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OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                          linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
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MOLECULE TYPE:
US-08-513-9748-129
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LOCATION: 1
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                                                                                                                                                        0; Indels
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APPLICANT: Hosoya, Masaki
APPLICANT: Hosoya, Masaki
APPLICANT: Fujii, Ryo
APPLICANT: Ohtaki, Tetsuya
APPLICANT: Pukusumii, Shoji
APPLICANT: Pukusumii, Shoji
APPLICANT: Ohgi, Kazuhiro
TITLE OF INVENTION: G PROTEIN COUPLED RECEPTOR PROTEIN,
TITLE OF INVENTION: PRODUCTION, AND USE THEREOF
CORRESPONDENCES: 380
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: DIKE, BRONSTEIN, ROBERTS & CUSHMAN, STREET: 130 Water Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                          Score 15; DB 3; Pred. No. 15; 0; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/JP95/01599
FILING DATE: 10-AUG-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 7-093989
FILING DATE: 19-AUG-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 7-057186
FILING DATE: 16-MAR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 7-007177
FILING DATE: 20-JAN-1995
PRIOR APPLICATION NUMBER: JP 6-326611
FILING DATE: 28-DEC-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 6-326611
FILING DATE: 28-DEC-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/513,974B
FILING DATE: 14-SEP-1995
CLASSIFICATION: 536
             OTHER INFORMATION: antisense sequence US-09-377-310-35
                                                                                                                                                                                                                                                                                                                       US-08-513-974B-129
US-08-513-974B-129
Sequence 129, Application US/08513974B
Patent No. 6114139
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
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APPLICATION NUMBER: JP 6-236356
FILING DATE: 30-SEP-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: JP 6-270017
FILING DATE: 02-NOV-1994
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: JP 6-236357
FILING DATE: 30-SEP-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: JP 6-189274
FILING DATE: 11-AUG-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 6-189273
                                                                                                          75.0%;
                                                                                                     Query Match
Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                                                      3 ttggtttccaatcgg 17
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MEDIUM TYPE: Floppy
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FEATURE:
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Gaps

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TITLE OF INVENTION: GRB3-3 GENE, VARIANTS AND USES THEREOF NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/612,857
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 11.8; DB 2;
Pred. No. 8.2e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3: Swanson & Bratschun, L.L.C.
8400 E. Prentice Avenue, Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATKINSON, BRENT
FENTION: SYSTEMATIC EVOLUTION OF
FENTION: LIGANDS BY EXPONENTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           other nucleic acid
/desc = "oligonucleotide IV"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: FR 93-10971
FILING DATE: 15-SEP-1993
PROR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/FR94/00542
FILING DATE: 09-MAY-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TALGEBOOK (610) 454-3809

TELEFAX: (610) 454-3809
                                                                                                               Rhone-Poulenc Rorer Inc.
                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 29, Application US/08612895A Patent No. 5763177 GENERAL INFORMATION:
APPLICANT: GOLD, LARRY APPLICANT: WILLIS, MICHAEL
                                                                                                               ADDRESSEE: Rhone-Poulenc Ro
STREET: 500 Arcola Rd. 3C43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: KOCH, TAD APPLICANT: RINGQUIST, STEVEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                59.0%;
86.7%;
                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: JENSEN, KIRK
APPLICANT: ATKINSON, BREN
TITLE OF INVENTION: LIGAN
TITLE OF INVENTION: ENRIC
TITLE OF INVENTION: OF NU
TITLE OF INVENTION: OF NU
TITLE OF INVENTION: SOLUT
  TOCQUE, Bruno
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (610)454-38.
TELEFAX: (610)454-3808
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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Matches 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 TCGTTTCCAAACGGA 16
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Swanson &
                                                                                                                                                                    Collegeville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
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                                                                                                                                                                                                                               USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; MOLECULE TYPE:
; DESCRIPTION:
US-08-612-857-9
                                                                                                                                                                                                                       COUNTRY: US
ZIP: 19426
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
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  APPLICANT:
                                                                                                                                                                       CITY: (STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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/note= "synthetic 5' primer used to create
recognition sites"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         A method for production of proteins in
                                                                                                                                                                                                                                                                      RESULT 5
US-08-948-591-8/C
i Sequence 8, Application US/08948591
; Patent No. 5939287
; GENERAL INFORMATION:
APPLICANT:
TITLE OF INVENTION: A method for production of proteins in TITLE OF INVENTION: yeast
NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
ADDRESSEE: Blich, Stewart, Kolasch and Birch
STREET: P.O. Box 747
; CTY: Falls Church
STRATE: VA
COUNTRY: USA
COUNTRY: USA
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: TEM PC COMPATIBLE
COMPUTER: PLOPSYMS-DOS
SOFTWARE: PATENTION DATA:
CURRENT APPLICATION DATA:
FILING DARF:
FILING DAFF:
FI
Score 12; DB 1; Lengtn ou, Pred. No. 6.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               60.0%; Score 12; DB 2; Length 3U; 75.0%; Pred. No. 6.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature
LOCATION: 5.10
OTHER INFORMATION: /function= "Kpn1 restriction site"
                       Query Match 60.0%; Score 12; DB Best Local Similarity 75.0%; Pred. No. 6.6e Matches 15; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PILING DATE:
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/295,676
FILING DATE: 09-SEPT-1994

FILING DATE: 09-SEPT-1994
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Sequence 9, Application US/08612857
Patent No. 5811048
GENERAL INFORMATION:
APPLICANT: SCHWEIGHOFFER, Fabien
                                                                                                                                       1 tgttggtttccaatcggacc 20
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: LOCATION: 1..30
: OTHER INFORMATION: OTHER INFORMATION: :
: OTHER INFORMATION: US-08-948-591-8
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Best Local Similarity
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ADDRESSEE: Swanson &
STREET: 8400 E. Prent
    MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCT-US94-10562A-29
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US-09-093-293-29
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Pred. No. 8.7e+02;
7; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: SYSTEMATIC EVOLUTION OF TITLE OF INVENTION: LIGANDS BY EXPONENTIAL TITLE OF INVENTION: ENRICHMENT: PHOTOSELECTION TITLE OF INVENTION: OF NUCLEIC ACID LIGANDS AND TITLE OF INVENTION: SOLUTION SELEX NUMBER OF SEQUENCES: 64
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
CADDRESSEE: SWANSON & BRATSCHUN, L.L.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 29, Application US/09093293
Patent No. 6001577
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: GOLD, LARRY
APPLICANT: WILLIS, MICHAEL
APPLICANT: WILLIS, MICHAEL
APPLICANT: STEVEN
APPLICANT: JENSEN, KIRK
APPLICANT: ATRINSON, BRENT
APPLICANT: ATRINSON, BRENT
TITLE OF INVENTION: LIGANDS BY EXPONENTIAL
                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.44 MG
MEDIUM TYPE: storage
COMPUTER: IBM
OPERATING SYSTEM: MS-DOS
SOFTWARE: WordPerfect 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/612,895A
FILING DATE: 03 August 1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Swanson & Bratschun, L.L.C. STREET: 8400 E. Prentice Avenue, Suite 200 CITY: Englawood STATE: Colorado
                                                                                                                                                                                                                                                                                         CASSIFICATION DATA:

APPLICATION DATA:

APPLICATION DATA:

APPLICATION NUMBER: PCT/US94/10542

FILING DATE: 18 September 1994

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/123,935

FILING DATE: 17 September 1993

APPLICATION NUMBER: 08/143,564

FILING DATE: 17 September 1993

APPLICATION NUMBER: 08/143,564

FILING DATE: 25 October 1993

ATORNEY AGENT INFORMATION:

NAME: BATTY J. SWARSON

REGISTRATION NUMBER: 33,215

REFERENCE/DOCKET NUMBER: NEXIO/PCT-US

TELECOMMUNICATION INFORMATION:

TELEFONE: (303) 793-333

INFORMATION FOR SEQ ID NO: 29:

SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 80111
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.44 MG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 59.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 59.0
Best Local Similarity 40.0
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 35 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 tgttggtttccaatc 15
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18 UGUUUGUUUCCACUC 32
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Colorado
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US-08-612-895A-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-093-293-29
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APPLICANT: WOLLIS, MICHAEL
APPLICANT: ROGH, TAD
APPLICANT: RINGQUIST, STEVEN
APPLICANT: JENSEN, RIRK
APPLICANT: ATKINSON, BRENT
TITLE OF INVENTION: SYSTEMATIC EVOLUTION OF
TITLE OF INVENTION: LIGANDS BY EXPONENTIAL
TITLE OF INVENTION: OF NUCLEIC ACID LIGANDS AND
TITLE OF INVENTION: SOLUTION SELEX
NUMBER OF SEQUENCES: 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3: Swanson & Bratschun, L.L.C.
8400 E. Prentice Avenue, Suite 200
                                                                                                PRILING DATE:

"CARASIEICATION DATA:

"FILING DATE:
"CALASSIEICATION DATA:

PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/612,895

FILING DATE: 03 August 1996

PRIOR APPLICATION NUMBER: PCT/US94/10542

FILING DATE: 18 September 1994

PRIOR APPLICATION NUMBER: PCT/US94/10542

FILING DATE: 18 September 1993

PRIOR APPLICATION NUMBER: 08/13,564

FILING DATE: 25 October 1993

APPLICATION NUMBER: 07/714,131

FILING DATE: 10 JUNE 1991

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 07/714,131

FILING DATE: 10 JUNE 1991

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 07/931,473

FILING DATE: 11 JUNE 1990

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 07/931,473

FILING DATE: 17 AGGUST 1992

ATTORNEY/AGENT INFORMATION:

NAME: BAILY J. SWARSON

REGISTRATION NUMBER: NEX10/US-CON

TELECOMMUNICATION INDREMATION:

NAME: BAILY J. SWARSON

RESISTRATION NUMBER: NEX10/US-CON

TELECOMMUNICATION INDREMATION:

TELECOMMUNICATION INDREMATION:

TELECOMMUNICATION FOR SEQ ID NO: 29:

SEDERICE CHARATTERISTICS:

SEDERICE CHARATTERISTICS:

SEDERICE CHARATTERISTICS:

SEDERICE CALCASTANTION:

SEDERICE CHARATTERISTICS:

SEDERICE CHARATTERISTICS:
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APPLICANT: GOLD, LARRY APPLICANT: WILLIS, MICHAEL
COMPUTER: IBM OPERATING SYSTEM: MS-DOS SOFTWARE: WORDPEFFECE 8.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ouery Match
Best Local Similarity 40.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 35 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 tgttggtttccaatc 15
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Englewood

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Gaps
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                                                                                                                                                                                                                                                                                                                                                                     STREET: 2000 Pennsylvania Avenue, NW, suite 5500 CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/765,783A
FILING DATE: 07-MAR-1997
CLASSIFICATION: 530
PRIOR APPLICATION NUMBER:
FILING APPLICATION NUMBER:
APPLICATION NUMBER:
ATTORNEY-AGENT INFORMATION:
NAME: MUTAShige, Kate H
REGISTRATION NUMBER: 29,959
REFERENCE/POCKET NUMBER: 35029-20001.20
TELEPOMMUNICATION INFORMATION:
TELEPHONE: 202-887-1500
                                                                                                                                                                                             APPLICANT: Sato, Koh
APPLICANT: Tsuchiya, Masayuki
APPLICANT: Yanazaki, Tatsumi
TITLE OF INVENTION: Reshaped Human Antibody
TITLE OF INVENTION: Interleukin-8
NUMBER OF SEQUENCES: 105
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COCATION: 1...34
CTHER INFORMATION: LTW1 sequence US-08-765-783A-42
                                                           Sequence 42. Application US/08765783A
Patent No. 5994524
GENERAL INFORMATION:
APPLICANT: Matsushima, Kouji
APPLICANT: Matsumoto, Yoshihiro
APPLICANT: Yamada, Yoshiki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-921-100-42/C
Sequence 42, Application US/08921100
Patent No. 6024956
GENERAL INFORMATION:
APPLICANT: MATSUSHIMA, KOUJI
APPLICANT: AMATSUSHIMA, ROSHIHIRO
APPLICANT: YAMADA, YOSHIKI
                                                                                                                                                                                                                                                                                                                                                        MORRISON & FOERSTER
                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: DC
COUNTRY: USA
ZIP: 20006-1888
COMPUTER READABLE FORM:
MEDIUM TYPE: D1sAvette
COMPUTER: IBM COMPALIDIE
OPERATING SYSTEM: DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3 ttggtttccaatcggacc 20
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SEQUENCE CHARACTERISTICS:
LENGTH: 34 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 77.8
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 8, Application US/09163444

Sequence 8, Application US/09163444

Sequence 8, Application US/09163444

Patent No. 6251596

Patent No. 6251596

PAPLICANT: Cook, James W.

APPLICANT: Cook, James W.

APPLICANT: Bulawa, Christine

TITLE OF INVENTION: ASPERGILLUS N-MYRISTOYL TRANSFERASE GENES AND POLYPEPTIDES

TITLE OF INVENTION: AND USES THEREOF

FILE REFERENCE: 07334/089001

CURREWY APPLICATION NUMBER: US/09/163,444

CURREWY FILING DATE: 1998-09-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ő
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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Pred. No. 8.8e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 11.8; DB 5;
Pred. No. 8.7e+02;
                                                                                  MEDIUM TYPE: Diskette, 3 1/2 inch, 1.44 MB MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; ORGANISM: Aspergillus fumigatus N-myristoyl
US-09-163-444-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 10
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                        COMPUTER: IBM
OPERATING SYSTEM: MS-DOS
SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
FILING DATE: 16 SEPTEMBER 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NEX10/PCT
                                                                                                                                                                                                                                                                          PRICE ASSISTANCE OF THE APPLICATION DATA:

APPLICATION NUMBER: 08/123,935
FILING DATE: 17 September 1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/143,564
FILING DATE: 25 October 1993
ATTORNEY/AGENT INFORMATION:
NAME: BAITY J. SWANSON
REFERENCE/DOCKET NUMBER: 33,215
REFERENCE/DOCKET NUMBER: NEXIO/PC
TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 793-3333
TELEFAX: (303) 793-333
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                59.0%;
86.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    59.0%;
40.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 59.0
Best Local Similarity 40.0
Matches 6; Conservative
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Best Local Similarity 86.7
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 35 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 tgttggtttccaatc 15
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18 UGUUUGUUUCCACUC 32
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                                                           COMPUTER READABLE FORM:
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Colorado
                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TOPOLOGY: ; TOPOLOGY: ; PCT-US94-10562A-29
                                      80111
                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 8
LENGTH: 39
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Pred. No. 1.1e+03;
0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/902,201
FILING DATE: 29-JUL-1997
CLASSIFICATION 1424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/437,328
FILING DATE: 09-MAY-1995
ATTONNEY/AGENT INFORMATION:
ANALE: WITH STREET STR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: MATSUSHIMA, KOUJI
APPLICANT: MATSUNTO, YOSHIHIRO
APPLICANT: YAMADA, YOSHIKI
APPLICANT: YAMADA, YOSHIKI
APPLICANT: YAMADA, YOSHIKI
APPLICANT: YOUNTAN, MASAYUKI
APPLICANT: YAMAZAKI, TATUMI
TITLE OF INVENTION: HUMANIZED ANTI-IL8 ANTIBODY
NUMBER OF SEGUENCES: 77
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
                                         Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   E: MORRISON & FOERSTER
2000 Pennsylvania Ave. NW, Suite 5500
                           CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/880,142
FILING DATE: 20-JUN-1997
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/345,145
FILING DATE: 28-NOV-1994
ATTORNEY/AGENT INFORMATION:
NAME: MUMBER: 29,959
REFERENCE/DOCKET NUMBER: 15580-0001.20
TELECOMMUNICATION INFORMATION:
TELEFHONE: (202) 887-1500
TELEFA: 90-4030
INFORMATION FOR SEQ ID NO: 42:
SEGUENCE CHARACTERISTICS:
LENGTH: 34 base pairs
INFORMANINGS: 151016
CTREET SECUENCE CHARACTERISTICS:
LENGTH: 34 base pairs
TELENGTH: 34 base pairs
TELENGTH: 34 base pairs
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REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 15580-0001.01
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 887-1500
          PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 42, Application US/08902201; Patent No. 6068840; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                58.0%;
77.8%;
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ZIP: 20006-1888
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3 ttggtttccaatcggacc 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               32 TTTGTTTCTAATGAGACC 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 58.0
Best Local Similarity 77.8
Matches 14; Conservative
          OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Washington
STATE: DC
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Pred. No. 1.1e+03;
0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                          COUNTRY: USA

ZIP: 2006-1888
COMPUTER READABLE FORM:
MEDIUW TYPE: FIDPPY disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN PC-DOS/MS-DOS
SOFTWARE: PATENTIN PC-DOS/MS-DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/921,100
FILING APPLICATION A14
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/437,323
FILING DATE: 09-MAY-1995
ATTORNEY/AGENT INFORMATION:
NAME: MURASHIGE, KATE H.
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 15580-0001.02
TELEPHONE: (202) 887-1500
TELEPHONE: (202) 887-1500
TELEPHONE: (202) 887-1003
TELEPHONE: (202) 887-1053
TELEPHONE: CANADATION:
TELEPHONE: CANADATION
SEQUENCE CHARACTERISTICS:
LEMOTH: A14 base pairs
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APPLICANT: MATSUMOTO, YOSHIRIO
APPLICANT: SAMADA, YOSHIKI
APPLICANT: SATO, KOH
APPLICANT: TSUCHIYA, MASSAYUKI
APPLICANT: TSUCHIYA, MASSAYUKI
APPLICANT: TAMAZAKI, TATUMI
TITLE OF INVENTION: HUMANIZED ANTI-IL8 ANTIBODY
NUMBER OF SEQUENCES: 77
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
                                                                                                                                                                                                       ADDRESSEE: MORRISON & FOERSTER
STREET: 2000 Pennsylvania Ave. NW, Suite 5500
                                                              APPLICANT: YAMAZAKI, TATUMI
TITLE OF INVENTION: HUMANIZED ANTI-IL8 ANTIBODY
NUMBER OF SEQUENCES: 77
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: MORRISON & FOERSTER
STREET: 2000 Pennsylvania Ave. NW, Suite 5500
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 42, Application US/08880142
Patent No. 6048972
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
ZIP: 20006-1888
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             58.0%;
77.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 77.8
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
                                                                                                                                                                                                                                      STREET: 2000 Penr
CITY: Washington
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US-08-880-142-42/c
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-921-100-42
                                                                                                                                                                                                                                                                                                            STATE:
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                                                                                                                                                                                                                                   Score 11.6; DB 3; Length 34;
Pred. No. 1.1e+03;
0; Mismatches 4; Indels
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2000 Pennsylvania Avenue, NW, suite 5500
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 42, Application US/09416557
Patent No. 6245894
GENERAL INFORMATION:
APPLICANT: Matsushima, Kouji
APPLICANT: Matsushima, Kouji
APPLICANT: Yamada, Yoshiki
APPLICANT: Yamada, Yoshiki
APPLICANT: Tatchiya, Masayuki
APPLICANT: Tatchiya, Masayuki
TITLE OF INVENTION: Reshaped Human Antibody to
TITLE OF INVENTION: Interleukin-8
NUMBER OF SEQUENCES: 105
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZOUNTY SOURCE TO SOURCE SOURCE SOURCE TO SOURCE TO SOURCE SOURCE FASTER DOS SOFTWARE: FASTER OF WINDOWS VERSION 2.0 CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/416,557
FILING DATE: 12-October-1999
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/765,783
FILING DATE: 7-March-1997
ATTORNEY,AGENT INFORMATION:
NAME: Murashige, Kate H
REGISTRATION NUMBER: 29,959
REFRENCE/FOOCKET NUMBER: 35029-20001.10
TELEPHONE: 202-887-100
TELEFAX: 202-887-100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; NAME/KEY: Other
; LOCATION: 1...34
; OTHER INFORMATION: LTW1 sequence
US-09-416-557-42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   42:
| TELEFAX: (202) 887-0763
| TELEX: 90-4030
| INFORMATION FOR SEQ ID NO: 42:
| SEQUENCE CHARACTERISTICS:
| LENGTH: 34 Dase pairs
| TYPE: nucleic acid
| STRANDEDNESS: single
| JOPOLOGY: linear
| US-08-902-201-42
                                                                                                                                                                                                                                   Query Match 58.0%;
Best Local Similarity 77.8%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 34 base pairs
LYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                          32 TTTGTTTCTAATGAGACC 15
                                                                                                                                                                                                                                                                                                                      3 ttggtttccaatcggacc 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Washington
STATE: DC
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 15
US-09-416-557-42/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE:
STREET: 20
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Search time 876.95 Seconds (without alignments)
14.320 Million cell updates/sec
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| SIDS8/gcgdata/geneseq/geneseqn/NA1980.DAT:*
| SIDS8/gcgdata/geneseq/geneseqn/NA1981.DAT:*
| SIDS8/gcgdata/geneseqn/geneseqn/NA1981.DAT:*
| SIDS8/gcgdata/geneseqn/geneseqn/NA1991.DAT:*
| SIDS8/gcgdata/geneseqn/geneseqn/geneseqn/NA1991.DAT:*
| SIDS8/gcgdata/geneseqn/geneseqn/NA1991.DAT:*
| SIDS8/gcgdata/geneseqn/geneseqn/NA1991.DAT:*
| SIDS8/gcgdata/geneseqn/geneseqn/RA1991.DAT:*

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GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             730101 seqs, 313950809 residues
                                                                                                                                                                                                                                                                                                  October 2, 2001, 16:18:42;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

    nucleic search, using sw model

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20
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Human focal adhesi Human focal adhesi HPV E6/7 region pr Cellulomonas fimi Description AAA12990 AAZ54999 AAQ04950 AAC65547 AAQ48576 a 22 22 22 11 22 12 20 12 16 Match Length DB 20 119 119 117 121 121 120 120 75.0 65.0 63.0 63.0 63.0 62.0 62.0 62.0 62.0 Query Score 12.6 12.6 12.6 12.4 12.2 12.2

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	20 AAV82U04 14 AAQ48878 18 AAT60126 21 AAC55244 19 AAV50662	20 AA209476 21 AA69516 22 AA655629 22 AA63216 21 AA72212 21 AA72212 22 AA722349 22 AA723349 22 AA723349 22 AA723349 21 AAC6475 21 AAC6476 21 AAC6606 21 AAC6606 22 AA71952 23 AAC6606 24 AAC6606 25 AAC6606 26 AAC6606 27 AAC6606	21 AAA65067 18 AAV79498 20 AAX85630 20 AAV91187 20 AAZ03515
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22 61. 22 61. 22 61. 22 60.	00000	11.8 8 59.0 11.1 8 8 59.0 11.1 8 8 59.0 11.1 8 8 59.0 11.1 8 8 59.0 11.1 8 8 59.0 11.1 6 58.0 11.1 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6	1.6 58. 1.6 58. 1.6 58. 1.4 57.
c 12 c 14 c 14 c 15 c 16	14444	00 000 0 00 0 42022000000000000000000000	c 41 42 43 44 45

ALIGNMENTS

Human; focal adhesion kinase; FAK; signal transduction; cancer; embryonic development disorder; angiogenic disorder; wound healing; antisense; phosphorothioate; ss. Human focal adhesion kinase antisense sequence #13 BP. 99US-0377310. 99US-0377310. AAC65547 standard; DNA; 20 (first entry) (ISIS-) ISIS PHARM INC. Monia BP, Gaarde WA; Homo sapiens 19-AUG-1999; 19-AUG-1999; US6133031-A 17-OCT-2000 12-FEB-2001 AAC65547; AAC65547 RESULT

New antisense compounds for inhibiting focal adhesion kinase expression, especially useful for inhibiting retinal neovascularization, or for diagnosing and treating e.g. colon cancer

WPI; 2001-006141/01.

Oligonucleotide ca Human genomic SNP Nucleotide sequenc Arabidopsis Ačxla Human B-raf substr Primer A (Group 11

AAA36501 AAZ99639 AAC66935 AAV93672 AAQ95843

Neisseria species

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The probe is used to detect benign and/or malignant human papilloma virus. The probe binds to the \rm E6 and/or \rm E7 region of the virus.
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xylo-oligosaccharide production; chimeric xylanase; PCR primer; ss.
                                                                                                                                                                                                                                                                                 Human papilloma virus; HPV; E6; E7; benign; malignant; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detecting benign and/or malignant human papilloma virus - by detecting DNA sequence of E6 and/or E7 region of human papilloma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cellulomonas fimi xylanase PCR primer, SEQ ID NO:6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   65.0%; Score 13; DB 14; 1
100.0%; Pred. No. 7.6e+02;
tive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 17; 18pp; Japanese.
                                                                                                                                  BP.
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                                                                                          RESULT 3
AAQ48576/c
ID AAQ48576 standard; DNA; 19
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Best Local Similarity 100.
Matches 13; Conservative
                        ttggtttccaatcgg 17
                                                                                                                                                                                                                                            HPV E6/7 region probe.
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14 TGTTGGTTTCCAA 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1993-277497/35
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                                                                                                                                                                                                         22-FEB-1994
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                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                     AAQ48576;
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ID AAA1
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                                  The present invention describes a number of phosphorothioate antisense sequences to the human focal adhesion kinase (FAK) protein. This protein is involved in integrin-mediated signal transduction, and is implicated in cancer, particularly colon, breast and oral tumours, embryonic development disorders, anglogenic disorders and wound healing. The antisense sequences, including the one shown here, can be used in the treatment of all of these.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes a number of phosphorothioate antisense sequences to the human focal adhesion kinase (FAK) protein. This protein is involved in integrin-mediated signal transduction, and is implicated in cancer, particularly colon, breast and oral tumours, embryonic development disorders, angiogenic disorders and wound healing. The antisense sequences, including the one shown here, can be used in the treatment of all of these.
                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New antisense compounds for inhibiting focal adhesion kinase expression, especially useful for inhibiting retinal neovascularization, or for diagnosing and treating e.g. colon cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; focal adhesion kinase; FAK; signal transduction; cancer; embryonic development disorder; angiogenic disorder; wound healing; antisense; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                  ö
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                                                                                                                                                                                                                                          100.0%; Score 20; DB 22; Length 20; 100.0%; Pred. No. 0.16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human focal adhesion kinase antisense sequence #33.
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                                                                                                                                                                                      Sequence 20 BP; 3 A; 5 C; 5 G; 7 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15 BP; 2 A; 3 C; 4 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 2
                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 75.0%; Score 15; DB Best Local Similarity 100.0%; Pred. No. 66; Matches 15; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 2; Column 25; 30pp; English.
Claim 15; Column 23; 30pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                          Query Match 100.
Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Monia BP, Gaarde WA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-006141/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-FEB-2001
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Gaps

; 0

Length 19; Indels

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AAZ53015 to AAZ54536, AAZ54577 to AAZ54615, and AAY74253 to AAX75941 represent novel Neisseria meningitis and N. gonorrheae polynucleotides and polypeptides. AAZ5457 to AAZ54576 and AAZ54616 to AAZ55473 represent PCR primers used in the exemplification of the present invention. The polypeptides, the polynucleotides, antibodies and compositions of the invention can be used as vaccines, as diagnostic reagents, and as immunogenic compositions. The polypeptides can be used in the manufacture of medicaments for treating or preventing infection due to Neisserial bacteria (e.g. meningitis and septicaemia), to detect the presence of Neisseria bacteria, or to raise antibodies. They may also be used to screen for agonists or antagonists, which may themselves may also be used in gene therapy protocols.
                                                                                                                                  Masignani V, Mora M;
Scalato E, Scarselli M;
                                                                                                                                                                                                                                                             Novel Neisserial polypeptides predicted to be useful antigens for vaccines and diagnostics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 35;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         63.0%; Score 12.6; DB 21;
78.9%; Pred. No. 1.38+03;
ive 0; Mismatches 4;
                                                                                                                               Fraser C, Galeotti C, Grandi G, Hickey E,
Petersen J, Pizza M, Rappuoli R, Ratti G,
Tettelin H, Venter JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 35 BP; 18 A; 5 C; 7 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                          Example 16; Page 150; 1453pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
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  98US-0103796.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ04950 standard; DNA; 43
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Best Local Similarity 78.9
Matches 15; Conservative
                                                                                       (GENO-) INST GENOMIC RES.
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                                                                                                                                                                                                                  WPI; 2000-062150/05.
                                                                   (CHIR ) CHIRON CORP
  09-OCT-1998;
25-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-NOV-1989;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates a novel chimeric xylanase (AAY81494) and to DNA encoding it (AAA12985). The chimeric xylanase comprises residues 1-244 of Streptomyces olivaceovitidis xylanase and residues 244-354 of Cellulomonas fini xylanase. Xylanase and residues 244-354 of Cellulomonas fini xylanase. Xylanase and cesidues 244-354 of Cellulomonas fini xylanase. Xylanase also known as endo-1.4-bera-xylan acomponent of the hemicallulose in plant cell walls, into xylo-oligosaccharides and xylose. Xylanase is used in a range of industrial processes. It is used to produce xylo-oligosaccharides from xylan from broad-leafed trees, and is used in wood- pulp bleaching to reduce the amount of chlorine required for this process. Xylo-oligosaccharides can be used as ingredients in foodstuffs and as water-retaining material in cosmetics. The chimeric xylanase of the invention does not generate xylose monomers during the hydrolysis of xylan. It is therefore useful for efficient and reliable xylo-oligosaccharide production. Sequences AAA12990-A1291 represent PCR primers used in an exemplification of the present invention to amplify a primers used in an exemplification of the present invention to amplify a facilian process.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                    Modified xylanase gene for use in foodstuff industry, has base sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Neisseria meningitidis; Neisseria gonorrheae; antigen; vaccine; antigenic; diagnosis; immunogenic; infection; meningitis; septicaemia; antibacterial; gene therapy; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       244-354 of the protein. The PCR product was then used to generate DNA encoding the chimeric xylanase via a second round of PCR using primers AAA12987 and AAA12991.
                                        (NORQ ) NORIN-SUISANSHO SHOKUHIN SOGO KENKYUSHOCHO.
(SEIB-) SEIBUTSUKEI TOKUTEI SANGYO GIJUTSU KENKYU SUISHIN KIKO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               63.0%; Score 12.6; DB 21; 78.9%; Pred. No. 1.3e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Neisseria species ORF cloning PCR primer #384.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 35 BP; 3 A; 15 C; 9 G; 8 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                              Example 1; Page 10; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98US-0094869.
98US-0098994.
98US-0099062.
98US-0103749.
  99JP-0071715
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2 gttggtttccaatcggacc 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98US-0083758
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99WO-US09346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ54999 standard; DNA; 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-MAR-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15; Conservative
                                                                                                                                                                        of 1065 nucleotides
                                                                                                         WPI; 2000-306509/27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
Matches 15; Conserv
17-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Neisseria sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-SEP-1998;
09-OCT-1998;
09-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            W09957280-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-NOV-11999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              31-JUL-1998
02-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ54999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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RESULT 5

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sequences used in the exemplification of the present invention. AAA35948 to AAA36632 represent nucleotide sequences containing SNPs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           χ.
                                                                                                                                                                                                                             AAZ99639 standard; DNA; 21 BP.
                                                                                 62.0%;
92.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       98EP-0116652.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99WO-EP06502
                                                                                                                                                                                                                                                                                 12-JUL-2000 (first entry)
                                         Sequence 17 BP; 5 A; 2 C;
                                                                          Query Match
Best Local Similarity 92.9
Matches 13; Conservative
                                                                                                                                    5 ggtttccaatcgga 18
                                                                                                                                                            1 ggtttccaattgga 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lipford GB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-256970/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200014217-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-MAR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Wagner H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                       AAZ99639;
                                                                                                                                                                                                               AAZ99639
    SSXS
                                                                                                                                                              a
                                                                                                                                     ò
                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A method has been developed for detecting the presence or absence of a single nucleotide polymorphism (SNP) allele in a genomic sample. The method comprises preparing a reduced complexity genome (RCG) from the genomic sample and analysing the RCG for the presence or absence of a SNP allele. The method can be used to characterise a tumour, to generate a genomic pattern for an individual genome or to generate a genomic classification code for a genome. The method can be used to assess whether a subject is at risk for developing a disease or to identify a set of SNP alleles associated with a disease. The method can also be used to perform linkage analysis. AAA35944 to AAA35947 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Detection of single nucleotide polymorphisms in genomes by preparation and analysis of reduced complexity genomes, useful for genotyping, fingerprinting and determining allele frequency of SNPs -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; single nucleotide polymorphism; SNP; genotyping; DNA analysis;
allels specific ollgonucleotide, ASO; reduced complexity genome; RCG;
genomic classification; identification; DNA fingerprinting;
tumour characterisation; hybridisation; ss.
                                                                                                                                                                                                                                                       Gaps
                                                                                         Factor VIII analogue is sufficiently mutated from the original gene that it is not recognised by blood Abs of the patient. The analogue is composed of human FVIII but carries the B-domain of human FV in place of the FVIII B domain.

Oligonucleotides can be used to alter the profile of the blood factor without significantly altering its activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human genomic SNP allele specific oligonucleotide SEQ ID NO:566.
                                                                                                                                                                                                                                                       .
0
                                                                                                                                                                                                                              Length 43;
                                                                                                                                                                                                                                                       Indels
                          New hybrid DNA encoding hybrid procoagulant proteins prepd. by modifying DNA encoding human factor VIII.
                                                                                                                                                                                                                          Score 12.6; DB 11;
Pred. No. 1.4e+03;
); Mismatches 4;
                                                                                                                                                                                       Sequence 43 BP; 10 A; 10 C; 11 G; 12 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Charest A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MASI ) MASSACHUSETTS INST TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Housman DE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 70; 111pp; English.
                                                                                                                                                                                                                                                      ;
0
                                                                                                                                                                                                                                                                                                                                                                        AAA36501 standard; DNA; 17 BP
                                                                                                                                                                                                                             63.0%;
78.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98US-0101757
                                                                                                                                                                                                                                                                                1 tgttggtttccaatcggac 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99WO-US22283
                                                                                                                                                                                                                                                                                                         25 tgtaggatgccaatgggac 43
                                                                                                                                                                                                                                                                                                                                                                                                                           26-JUL-2000 (first entry)
                                                                  Disclosure; ; pp; English.
                                                                                                                                                                                                                            Query Match 63.0
Best Local Similarity 78.9
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Jordan B,
WPI; 1990-193265/25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-293181/25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200018960-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Landers JE,
                                                                                                                                                                                                                                                                                                                                                                                                  AAA36501;
                                                                                                                                                                                                                                                                                                                                                            AAA36501
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The present sequence represents a G-motif oligonucleotide of the invention. The specification describes compositions comprising G-motif oligonucleotides inhibiti activation of aligonucleotides. The G-motif oligonucleotides inhibit activation of artigen presenting cells by inhibiting the uptake of DNA by a cell, by stimulating natural killer cells, or by co-stimulating cytotoxic.

Tripophocytes. The G-motif oligonucleotides may be used for the productions of vaccines for treating septic shock, inflammation, autoimmune diseases (e.g. rheumatoid arthritis, Crohn's disease, sarcoidosis, multiple sclerosis, Kawasaki syndrome, graft-versus-host diseases (e.g. Streptococcal induced arthritis, Lyme arthritis, chronic inflammatory bowel disease, psoriasis vulgaris, experimental allergic encephalomyelitis, and insulin-dependent diabetes mellitus), bacterial infections, parasitic infections (e.g. Leishmaniasis or Toxoplasmosis), viral infections (e.g. Cytomegalovirus and human immunodeficiency virus)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           G-motif oligonucleotide; vaccine; Toxoplasmosis; viral infection; antigen presenting cell activation; natural killer cell; septic shock; cytocoxic T-lymphocyte; inflammation; autoimmune disease; rheumatoid arthritis; Crohn's disease; sarcoidosis; multiple sclerosis; Kawasaki syndrome; graft-versus-host disease; transplant rejection; helper T cell response 1-mediated disease; Lyme arthritis; streptococcal induced arthritis; chronic inflammatory bowel disease; psoriasis vulgaris; experimental allergic encephalomycalitis; insulin-dependent diabetes mellitus; bacterial infection; tumour; ss. parasitic infection; Leishmaniasis; spontaneous abortion; tumour; ss.
                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Compositions comprising G-motif oligonucleotides useful for treating e.g. septic shock, rheumatoid arthritis, diabetes and human
                                                                                                                                                                                                                                                          ö
                                                                                                                                                          Length 17;
                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleofide sequence of G-motif oligonucleotide PZ314.
                                                                                                                                                          Score 12.4; DB 21;
Pred. No. 1.6e+03;
5 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CPGI-) CPG IMMUNOPHARMACEUTICALS GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunodeficiency virus infections -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 34; 75pp; English
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c-raf; A-raf; B-raf; hammerhead ribozyme; hairpin ribozyme;

BP.

(first entry)

SSSX8

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target; substrate; catalyst; modulation; expression; Raf gene; delivery; screening; identification; synthesis; deprotection; purification; cancer; inflammation; psoriasis; non-hepatic ascites; infection; genetic drift; restenosis; rheumatoid arthritis; ss.
                                                                                                      Human B-raf substrate nucleotide position 2272.
                                   AAV93672 standard; RNA; 17
                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                       WO9850530-A2
                                                                                                                                                                                                                                                                                                                                         22-AUG-1997;
02-OCT-1997;
02-OCT-1997;
                                                                                                                                                                                                                                                                     05-MAY-1998;
                                                                               18-FEB-1999
                                                                                                                                                                                                                                              12-NOV-1998
                                                                                                                                                                                                                                                                                            19-DEC-1997;
                                                                                                                                                                                                                                                                                                                09-JUN-1997
03-JUL-1997
                                                                                                                                                                                                                                                                                                                                                                            05-NOV-1997
                                                                                                                                                                                                                                                                                                        09-MAY-1997
                                                        AAV93672;
             10
                         AAV93672
              RESULT
                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention provides a method for identifying genes encoding components of plant signal transduction pathways between mitochondrial function and metabolic status and nuclear gene expression. This involves the transformation of a plant with a vector encoding a reporter gene linked to an AOX promoter, mutagenesis of the plant to increase expression of the reporter gene and then determining the identity of the signal transduction gene of interest. This is useful in the production of transgenic plants with increased productivity, herbicide, stress and pest
(HIV)-infections), spontaneous abortions and tumours. They may also be used to induce proliferation of bone marrow cells, especially macrophage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Identifying genes encoding signal transduction components useful for producing transgenic plants, by transforming plants with vector encoding reporter gene, mutating the plant and identifying genes from
                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                   primer; ss.
                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24;
                                                                               Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                                      Indels
                                                                                                                                                                                                                                                                                                    Plant signal transduction pathway; reporter vector;
luciferase; herbicide resistance; pest resistance; PCR
                                                                               Score 12.4; DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 12.4; DB 22;
Pred. No. 1.6e+03;
0; Mismatches 1;
                                                                                           1.6e+03;
                                                                                                                                                                                                                                                                                Arabidopsis AOX1a gene promoter PCR primer #2.
                                           Sequence 21 BP; 2 A; 5 C; 3 G; 11 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 6 A; 5 C; 5 G; 8 T; 0 other;
                                                                                                      0; Mismatches
                                                                                           Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 54; 54pp; English.
                                                                                                                                                                                                             BP.
                                                                              62.0%;
92.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            62.0%;
ilarity 92.9%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                  99US-0136145
                                                                                                                                                                                                                                                                                                                                                                                                           22-MAY-2000; 2000WO-US14041
                                                                                                                                                                                                           AAC66935 standard; DNA; 24
                                                                                                                                                                                                                                                        (first entry)
                                                                                                      13; Conservative
                                                                                                                            1 tgttggtttccaat 14
                                                                                                                                                  21
                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UYNE-) UNIV NEBRASKA.
                                                                                                                                                                                                                                                                                                                                      Arabidopsis thaliana.
                                                                                                                                       WPI; 2001-025143/03.
                                                                                        Best Local Similarity
Matches 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 13; Conserv
                      precursor cells.
                                                                                                                                                                                                                                                                                                                                                              WO200071668-A2.
                                                                                                                                                                                                                                                         27-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                  20-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                     30-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rhoades DM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the mutant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      resistance
                                                                                                                                                                                                                                 AAC66935;
                                                                               Query Match
                                                                                                                                                  æ
                                                                                                                                                                                                AAC66935
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970S-0046059. 970S-0049002. 970S-0051718. 970S-0056808. 970S-0061321. 970S-0061324.

97US-0064866

98WO-US09249 97US-0068212

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comprises: (a) introducing into the system a random library of nucleic acid catalysts (NAC) having a substrate binding domain (SBD), comprises:

acid catalysts (NAC) having a substrate binding domain (SBD), comprising a random sequence, and a catalytic domain (CD); and (b) identifying NAC in systems where modulation has occurred and/or determining the sequence of at least part of the SBDs in such systems. Nucleic acid molecules with endonuclease activity and catalytic activity, from the present invention, are used to modulate gene expression in plant and mammalian cells and to cleave target nucleic acid, particularly for treating systemic diseases caused by specific RNA, e.g. cancer, inflammation, psoriasis, non-hepatic ascites and infection. They may also be used to determine craft and mutations in diseased cells and to determine craft RNA. Specifically NACS with RNA-cleaving activity that modulate expression of the Raf gene, are used to treat cancer, restenosis, with the level of c-raf. Introduction of sugar/phosphate modifications in character that the level of c-raf. Introduction of sugar/phosphate modifications associated mith the level of c-raf. Introduction of sugar/phosphate modifications associated madility against nuclease and activity. Any 90922 to ANY 93877 confinence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               method has been developed for the identification of a nucleic acid
                                                                                                                                                                                                                                                                                                                                                                    Identifying new catalytic nucleic acid that modulates selected processes - especially ribozymes that cleave Raf RNA for treating cancer, restenosis, and also new ribozymes and modified nucleoside
                                                                                                                                                                                           Workman CT;
                                                                               Burgin A, Jarvis r;
mic J, McSwiggen JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        triphosphates used as antiviral agents and synthons
                                                                                                                                                                                           Thompson J,
                                                                                         Beigelman L, Bellon L, Burgin
Kisich K, Matulic-Adamic J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                modulating the expression of a Raf gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 177; Page 172; 259pp; English.
                                                                                                                                                                                           Sweedler D,
(RIBO-) RIBOZYME PHARM INC.
                                                                                                                                          A, Kisich K,
Reynolds M, S
                                                                                                                                                                                                                                                                                WPI; 1999-009494/01.
                                                                                                                                               Karpeisky A,
                                                                                             Beaudry A,
                                                                                                                                                                                           Parry T,
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Sequence 17 BP; 5 A; 4 C; 4 G; 4 U; 0 other;

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Gaps

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Indels

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tggtttccaatcgg 17

4 11

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24

Query Match

AAQ95843;

AAQ95843

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PCR primers AAV23717-32 were used during the course of the invention. The specification describes Pseudomonas pseudoalcaligenes lipase variants. The variant lipases described in the specification has have better wash performance than the original lipase. It has improved removal of lipid stains, reduced calcium dependence, better compatibility with detergents or their components, increased hydrophobicity, altered substrate specificity and better one-cycle wash efficiency. The variant lipase is used in detergent and cleaning compositions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR primer; human; growth hormone; variant; cachexia; erythropoietin; erythrocyte formation; interferon; antiviral; antitumour; interleukin; immunomodulatory agent; colony-stimulating factor; proliferation stimulation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Variants of lipase from Pseudomonas containing specific amino acid substitutions - deletions or additions, having improved wash performance in detergent formulations
                                                                                                                                                                                                     Lipase; variant; improved wash performance; removal; lipid stain; reduced calcium dependence; one-cycle wash efficiency; detergent; cleaning composition; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 12.2; DB 19;
Pred. No. 2.1e+03;
0; Mismatches 3;
                                                                                                                                                               PCR primer used in the course of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 28 BP; 7 A; 7 C; 8 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Primer BB10 for human growth hormone gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 75; 104pp; English.
                                          BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP.
                                                                                                                                                                                                                                                                                                          Pseudomonas pseudoalcaligenes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61.0%;
82.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    96US-0029190.
96DK-0000902.
                                                                                                                                                                                                                                                                                                                                                                                                                              97WO-DK00345
                                          28
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                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3 ttggtttccaatcggac 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 82.4
nes 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (NOVO ) NOVO-NORDISK AS.
                                       AAV23724 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Okkels JS, Svendsen A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21 TIGGTITCGAACCGAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1998-230259/20.
                                                                                                                                                                                                                                                                                                                                              WO9808939-A1.
                                                                                                                     20-AUG-1998
                                                                                                                                                                                                                                                                                                                                                                                                                              26-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-AUG-1996;
27-AUG-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                      05-MAR-1998.
                                                                                                                                                                                                                                                                                   Synthetic
                                                                               AAV23724;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX21962;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Loca
Matches
  RESULT 12
                       AAV23724/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                automated genotyping. esp. fluorescence based. The primers correspond to the unique DNA sequence surrounding each marker, and PCR is used to detect each polymorphism. When the MMS show considerable polymorphism (ie. a difference in the number of repeats) between individuals, the markers can be particularly informative. The MMS can be ideal for linkage studies. Kits comprise at least 4 groups, of at least 3 sets, each comprising labelled primers for PCR amplification of the DNA. Group II primer pairs are shown in AAQ95841-82. The published size range of the D95180 allele is 220-265 bp, and the degree of heterozygosity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kit for automated genotyping contg. pairs of PCR primers - designed to amplify polymorphic nucleotide repeat sequences, arranged in sets each with a characteristic fluorescence label, useful e.g. in detection of disease related genetic rearrangement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The method aims to provide a collection of highly reproducible microsatellite marker sequences (MMS) at approx. 10-50 cM intervals throughout the human genome which can be detectably labelled. The MMS are polymorphic, simple sequence repeats and can be used in
                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
                                                                                                                                                                                                                                                                                                                                                                              primer; polymerase chain reaction; PCR; linkage study; locus; microsatellite marker sequence; automated genotyping; allele; polymorphism; detection; Homo sapiens; ss.
Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 20;
                                                                                                                                                                                                                                                                                                                                              Primer A (Group 11, set A) for marker D9S180, chromosome 9.
                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
Score 12.2; DB 20;
Pred. No. 2e+03;
                                       3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 12.2; DB 16;
Pred. No. 2e+03;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 5 A; 4 C; 6 G; 5 T; 0 other;
                                       4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Fig 7K-2; 104pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     population is about 63%
                                                                                                                                                                                                                      AAQ95843 standard; DNA; 20 BP
61.0%;
58.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94WO-US13945.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61.0%;
ilarity 82.4%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UYJO ) UNIV JOHNS HOPKINS.
                                                                                                                                                                                                                                                                                                      (first entry)
                                                                             2 gttggtttccaatcgga 18
                                                                                                                 1 gcugguuuccaaacaga 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    tggtttccaatcggacc 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4 tggtttggaatcgaacc 20
                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1995-215278/28.
                                                                                                     _ = ::: = ::
                     Similarity
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Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9515400-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-DEC-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                03-DEC-1993;
                                                                                                                                                                                                                                                                                                      21-FEB-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-JUN-1995
                     Best Local Sim
Matches 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Levitt RC;
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Gaps

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Best Loca Matches

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Length 28; Indels

Synthetic

Cox GN;

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The present invention concerns novel methods of producing proteins with free cysteine residues. The PCR primers AAA/A770-A74854 were used to amplify sequences encoding the human growth hormone, erythropoietin and IFN-alpha2 proteins, and signal sequences used to control their secretion, and change these sequences in order to alter their efficacy. One example of modifications is the addition of polyethylene glycol moieties. The proteins could then be produced in cell cultures and isolated for use in the treatment of disorders such as those associated with growth hormone, erythropoietin or IFN-alpha2.
                                                                                                                                                                                                                                                              Producing a soluble protein having a free cysteine comprises exposing a host cell expressing the soluble protein to a cysteine blocking agent, useful for producing recombinant human growth factor and erythropoietin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Producing a soluble protein having a free cysteine comprises exposing a host cell expressing the soluble protein to a cysteine blocking agent, useful for producing recombinant human growth factor and erythropoietin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Erythropoietin; growth hormone; IFN-alpha2; free cysteine residue; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 12.2; DB 21;
Pred. No. 2.1e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 32 BP; 7 A; 13 C; 6 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human growth hormone mutagenic primer BB33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rosendahl MS;
                                                                                                                                                                           Rosendahl MS;
                                                                                                                                                                                                                                                                                                                                                                      Example 2; Page 18; 86pp; English.
                                                                                                                                 (BOLD-) BOLDER BIOTECHNOLOGY INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (BOLD-) BOLDER BIOTECHNOLOGY INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             61.0%;
82.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-0116041.
                                                                                     99US-0116041.
                                         14-JAN-2000; 2000WO-US00931.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-JAN-2000; 2000WO-US00931
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA74783 standard; DNA; 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4 tggtttccaatcggacc 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cox GN, Doherty DH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Doherty DH,
                                                                                                                                                                                                                  WPI; 2000-476056/41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-476056/41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-JAN-1999;
                                                                                   14-JAN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer;
20-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA74783;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COX GN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA74783,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This sequence is a PCR primer for the human growth hormone gene. The invention relates to a variant of a member of the growth hormone (GH) superfamily that has a Cys residue: (a) replacing an amino acid (aa) in the loop regions, near the end of the alpha-helices or in front of, or after, the first amphipathic helix; (b) added to the N- or C- terminus; or (c) introduced between two as in the loop region, at the ends of the calpha-helices or in front of, or after, the first amphipathic helix. The variants are used for the same therapeutic purposes as the wild type sequences, e.g. for GH to treat short stature or cachexia; for sequences, e.g. for GH to treat short stature or cachexia; for crythropoietin (EPO) to stimulate erythrocyte formation; for interferons as antiviral, antitumour or immunomodulatory agents; for colony-stimulating factors or interleukins for stimulating proliferation, differentiation and function of particular types of (haematopoietic) colls. Incorporation of Cys at a non-essential position allows site-specific attachment of e.g. poly(ethylene glycol) (PEG) to create a conjugate with increased circulation time in vivo. This reduces the need for frequent injections and the cost of treatment. Modification with PEG also increases solubility and stability of the protein and reduces its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
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                                                                                                                                                                                                                                                                                                                                                                                           New variants of growth hormone superfamily proteins containing an extra cysteine - useful, e.g. for treatment of short stature or cachexia, to stimulate erythrocyte formation and as immunomodulator
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Erythropoietin; growth hormone; IFN-alpha2; free cysteine residue;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 12.2; DB 20;
Pred. No. 2.1e+03;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 32 BP; 7 A; 13 C; 6 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E. coli stII signal sequence PCR primer BB10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 16; 94pp; English.
                                                                                                                                                                                                                                                              (BOLD-) BOLDER BIOTECHNOLOGY INC.
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82.4%;
                                                                                                                                                                                                                  97US-0052516
                                                                                                                                                                        98WO-US14497
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAA7476
ID AAA74774 standard; DNA; 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-JAN-2001 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 61.0
Best Local Similarity 82.4
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                               WPI; 1999-132163/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Escherichia coli.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200042175-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  immunogenicity
                                           Homo sapiens.
                                                                                 WO9903887-A1
                                                                                                                                                                                                                  14-JUL-1997;
                                                                                                                                                                      13-JUL-1998;
                                                                                                                          28-JAN-1999
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AAA74774;

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The present invention concerns novel methods of producing proteins with free cysteine residues. The PCR primers AAA7470-A74854 were used to amplify sequences encoding the human growth hormone, errythropoietin and IFN-alpha2 proteins, and signal sequences used to control their secretion, and change these sequences in order to alter their efficacy. One example of modifications is the addition of polyethylene glycol modeties. The proteins could then be produced in cell cultures and isolated for use in the treatment of disorders such as those associated with growth hormone, erythropoietin or IFN-alpha2.
                                                    Example 4; Page 24; 86pp; English.
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; 0 0; Gaps Score 12.2; DB 21; Length 34; Pred. No. 2.2e+03; 0; Mismatches 3; Indels C 61.0%; Query Match
Best Local Similarity 82.4
Matches 14; Conservative

Sequence 34 BP; 8 A; 11 C; 9 G; 6 T; 0 other;

4 tggttccaatcggacc 20 ||| || || ||||||| || 29 TGGATTCTAATCGGATC 13 ò

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Search completed: October 2, 2001, 16:18:43 Job time: 15487 sec

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TYPE: DNA ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 20; Conservative
     us-09-377-310-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-377-310-32
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LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
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  0000000000
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Appl
                                                                                                                                                                                                                              (without alignments)
9.071 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                              ; Search time 417.38 Seconds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Description
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Sequence
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/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
                          GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                      - nucleic search, using sw model
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Match Length
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seq length: 50
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Sequence 2
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APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310B
CURRENT APPLICATION NUMBER: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
                        Sequence
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APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310B
CURRENT APPLICATION NUMBER: US/09/377,310B
NUMBER OF SEQ ID NOS: 1999-08-19
SOFTWARE: PatentIn Ver. 2.0
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US-09-219-932-17
US-08-243-542-20
US-08-477-407-20
US-08-377-1248-2219
US-08-373-1248-2219
US-08-373-1248-2019
US-08-373-1248-2019
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US-07-812-421-20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 12, Application US/09377310B
Patent No. 6133031
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 32, Application US/09377310B Patent No. 6133031 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                          APPLICANT: Draper, Kenneth
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
TITLE OF INVENTION: TRETHODS AND COMPOSITIONS FOR
TITLE OF INVENTION: TREATMENT OF RESTENOSIS AND
TITLE OF INVENTION: CANCER USING RIBOZYMES
NUMBER OF SEQUENCES: 2627
ADDRESSEE: Lyon & Lyon
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Pred. No. 79;
0; Mismatches 2;
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: 5.5 Cargae
COMPUTER: 18M Compatible
COMPUTER: 18M Compatible
COMPUTER: 18M Compatible
CORPUTER: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/435,628
FILING DATE: 0.5 MAY 1995
CLASSIFICATION S14
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/33,124
FILING DATE: January 13, 1995
APPLICATION NUMBER: 08/35,466
FILING DATE: MAY 18, 1994
APPLICATION NUMBER: 08/192,943
FILING DATE: PEDRUARY 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/987,132
FILING DATE: AUGUST 26, 1992
APPLICATION NUMBER: 07/987,132
FILING DATE: AUGUST 26, 1992
APPLICATION NUMBER: 07/986,422
FILING DATE: AUGUST 26, 1992
                                                                                                                                                                          ; Sequence 1431, Application US/08435628
; Patent No. 5817796
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 209/03
TELECOMUNICATION INFORMATION:
TELEPAX: (213) 955-0440
TELEX. (7-31) 955-0440
INFORMATION FOR SEQ ID NO: 1431:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                          APPLICANT: Stinchcomb, Dan T.
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Best Local Similarity 88.29
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TYPE: nucleic acid
STRANDEDNESS: single
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STATE: California
COUNTRY: U.S.A.
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                                                                                                            Length 15;
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                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Stinchcomb, Dan T.
APPLICANT: Draper, Kenneth
APPLICANT: Draper, Kenneth
APPLICANT: Drayer, James
APPLICANT: Jarvis, Thale
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR
TITLE OF INVENTION: TREATMENT OF RESTENOSIS AND
TITLE OF INVENTION: CANCER USING RIBOZYMES
CORRESPONDENCE ADDRESS:
                                                                                                            DB 3;
18;
                                                                                                          Score 15; DB
Pred. No. 18;
0; Mismatches
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88.2%; Pred. No. 79;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRILING DATE: May 18, 1994
APPLICATION NUMBER: 08/245,466
FILING DATE: May 18, 1994
APPLICATION NUMBER: 08/192,943
FILING DATE: PEDRUARY 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/936,422
FILING DATE: AUGUST 26, 1992
APTONNEY AGENT INFORMATION:
NAME: Warburg, Richard
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 209/035
TELEPHONE: (213) 489-1600
TELEPHONE: (213) 489-1600
TELEPKA: (213) 955-0440
TELERA: 67-3310
INFORMATION FOR SEQ ID NO: 1431:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRENDENDENSES: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/373,124A FILING DATE: January 13, 1995 PRIOR APPLICATION DATA:
; FEATURE:
; OTHER INFORMATION: antisense sequence
US-09-377-310-32
                                                                                                                                                                                                                                                                                                                     US-08-373-124A-1431/c
; Sequence 1431, Application US/08373124A
; Patent No. 5646042
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: 633 West Fifth Street STREET: 631 West Fifth Street STREET: Suite 4700
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                                                                                                          75.0%; ; 100.0%;
                                                                                                       Query Match 75.0
Best Local Similarity 100.
Matches 15; Conservative
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                                                                                                                                                                                                                    3 tgacatcagtagcat 17
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STATE: California
COUNTRY: U.S.A.
ZIP: 90071
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Best Local Similarity
Matches 15; Conserv
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Gaps
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GENERAL INFORMATION:
APPLICANT: Lonberg, Nils
APPLICANT: Kay, Kobert M.
TITLE OF INVENTION: Transgenic Non-Human Animals Capable of
TITLE OF INVENTION: Producing Heterologous Antibodies
NUMBER OF SEQUENCES: 77
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: William M. Smith
STREET: One Market Plaza, Steuart Tower, Suite 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
SOFTARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/834,539A
FILING DATE: 1992-02-05
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 13.8; DB 5;
                                                                                                                                                                     NAME: Osman, Richard A
REGISTRATION NUMBER: 35,627
REFERENCE/DOCKET NUMBER: FP-59232-PC/RAO
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred. No. 88;
0; Mismatches
                      APPLICATION NUMBER: PCT/US95/05265
                                                           CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/235,503
FILING DATE: 29-ARR-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30, 223
REFERENCE/DOCKET NUMBER: 1464
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-543-9600
TELEFAX: 415-543-5043
                                                                                                                                                                                                                                                     TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 64:
SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                         64:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  69.0%;
88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
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Best Local Similarity 88.2
Matches 15; Conservative
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 415-543-5043 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 42 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
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; MOLECULE TYPE:
PCT-US95-05265-64
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                                         FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
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GENERAL INCORMATION:
APPLICANT: TULATIK, INC.
TITLE OF INVENTION: TRANSCRIPTION FACTOR-DNA BINDING ASSAY
NUMBER OF SEQUENCES: 74
CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
STREET: San Francisco
CITY: San Francisco
STATE: California
                                 GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Peterson, Michael G
APPLICANT: Baichwal, Vijay R
APPLICANT: Strulovici, Berta
TTILE OF INVENTION: TRANSCRIPTION FACTOR-DNA ASSAY
NUMBER OF SEQUENCES: 75
CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
STREET: 4 Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1; Length 30;
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPPERTING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/235,503B
FILING DATE: 29-APR 1994
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
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                                                           Sequence 64, Application US/08235503B Patent No. 5563036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 29-ARR-1994
CLASSIFICATION: 435
ATGREEY-AGENT INFORMATION:
NAME: OSMAN, RICHARD A
REGISTRATION NUMBER: 36,627
REFERENCE/DOCKET NUMBER: A-59:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415, 398-3249
TELEY. 910 277299
INFORMATION FOR SEQ ID NO: 64:
SEQUENCE CHARACTERISTICS:
LENGTH: 30 Dasse pairs
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88.2%;
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Best Local Similarity 88.2
Matches 15; Conservative
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TOPOLOGY: linear
; MOLECULE TYPE: CDNA
US-08-235-503B-64
                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
ZIP: 94111-4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 94111-4187
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PCT-US95-05265-64/c
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5719032-Human Animals for
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Patent No. 5789550
GENERAL INFORMATION:
APPLICANT: Lonberg, Nils
APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transgenic No. 5789650-Human Anima
TITLE OF INVENTION: Producing Heterologous Antibodies
NUMBER OF SEQUENCES: 150
CORRESPONDENCES: Milliam M. Smith
STREET: One Market Plaza, Steuart Tower, Suite 2000
                                                                                                                                                                APPLICANT: Lonberg, Nils
APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transgenic No. 5719032-Human Anim
TITLE OF INVENTION: Producing Heterologous Antibodies
NUMBER OF SEQUENCES: 150
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWAREN PAINTED THE STATE OF THE SOFTWAREN PAPELICATION DATA:
APPLICATION NUMBER: US/08/645,641
FILING DATE: 20-MAY-1996
CLASSIFICATION: 800
PRIOR APPLICATION BOTH STATE:
APPLICATION DATA:
APPLICATION BOTH STATE:
APPLICATION BOTH STATE:
APPLICATION BOTH STATE:
APPLICATION BOTH STATE:
APPLICATION NUMBER: 03 / 1992
ATTONNEY/ACENT INFORMATION:
NAME: SMITH, WILLIAM M.
REGISTRATION NUMBER: 30, 223
REGISTRATION NUMBER: 14643-000913
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             68.0%; Score 13.6; DB 1;
80.0%; Pred. No. 1.2e+02;
ive 0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                   STREET: Two Embarcadero Center, 8th Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPATIBLE ISM PC COMPATI
                                                                     Sequence 64, Application US/08645641
Patent No. 5719032
GENERAL INFORMATION:
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TELEPHONE: 415-326-2400
TELEFRA: 415-326-2422
INFORMATION FOR SEQ ID NO: 64:
SEQUENCE CHARACTERISTICS:
LENGTH: 42 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 80.0
Matches 16; Conservative
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STRANDEDNESS: single
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California
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      RESULT 9
US-08-645-641-64/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Lonberg, Nils
APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transgenic No. 5661016-Human Animals for
TITLE OF INVENTION: Producing Heterologous Antibodies
NUMBER OF SEQUENCES: 197
CORRESPONDENCE ADDRESSE: Townsend and Townsend Khourie and Crew
STREET: One Market Plaza, Steuart Tower, Suite 200
                                                                                                 68.0%; Score 13.6; DB 1; Length 42; 80.0%; Pred. No. 1.2e+02; Live 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 42;
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Pred. No. 1.2e+02;
0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/990,860
FILING DATE: 16-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/810,279
FILING DATE: 17-DEC-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/853,408
FILING DATE: 18 WAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 14643-9-3
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/053,131
FILING DATE: 26-APR-1993
CLASSIFICATION: 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 64, Application US/08053131 Patent No. 5661016 GENERAL INFORMATION:
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ilarity 80.0%;
Conservative
                                                                                                 Query Match 68.0
Best Local Similarity 80.0
Matches 16; Conservative
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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STRANDEDNESS: single
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MEDIUM TYPE: Floppy
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California
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Best Local Similarity
Matches 16; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: US
US-07-834-539A-12
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ZIP: 94105 COMPUTER READABLE FORM:

94105

Length 42; Indels

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Sequence 12, Application US/08800353

Patent No. 5874299

GENERAL INFORMATION:
APPLICANT: Lonberg, Nils

APPLICANT: Any Robert M.
TITLE OF INVENTION: Transgenic No. 5874299-Human Animals Capable of TITLE OF INVENTION: Producing Heterologous Antibodies
NUMBER OF SEQUENCES: 77

CORRESPONDENCE ADDRESS:
ADDRESSEE: William M. Smith
STREET: One Market Plaza, Steuart Tower, Suite 2000
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 68.0%; Score 13.6; DB 1; Length 42; Best Local Similarity 80.0%; Pred. No. 1.2e+02; Matches 16; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 94105
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURREW APPLICATION DATA:
APPLICATION NUMBER: US/08/800,353
  APPLICATION NUMBER: US 07/810,279
FILING DATE: 17-DEC-1991
ATTORNEY/AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION NUMBER: 14643-9-4
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-326-2400
TELEPHONE: 415-326-242
INFORMATION FOR SEQ ID NO: 64:
SEQUENCE CHARACTERISTICS:
LENGTH: 42 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/834,539
FILING DATE: 1992-02-05
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 1464
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-543-9600
TELEFAX: 415-543-5043
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 12: SEQUENCE CHARACTERISTICS: LENGTH: 42 base pairs TYPE: nucleic acid STRANDENESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; MOLECULE TYPE: DNA (genomic) US-08-800-353-12
                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear MOLECULE TYPE: DNA (primer)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 cctgacatcagtagcatctc 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-08-800-353-12/c
                                                                                                                                                                                                                                                                                                                                                                                          US-08-096-762-64
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Fatent No. 5814318
GENERAL INFORMATION:
APPLICANT: Lonberg, Nils
TITLE OF INVENTION: Transgenic No. 5814318-Human Animals for TITLE OF INVENTION: Producing Heterologous Antibodies
NUMBER OF SEQUENCES: 210
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend Khourie and Crew
STREET: One Market Plaza, Steuart Tower, Suite 200
CITY: San Francisco
STRATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 94105
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/096,762
FILING DATE: 22-JUL-1993
CLASSIFICATION: 800
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFWWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
68.0%; Score 13.6; DB 1;
Best Local Similarity 80.0%; Pred. No. 1.2e+02;
Matches 16; Conservative 0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRICE APPLICATION: 800
PRICE APPLICATION DATA:
APPLICATION NUMBER: US 08/053,131
FILING DATE: 26-APR-1993
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
FILING DATE: 16-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/904,068
FILING DATE: 23-JUN-1992
PRIOR APPLICATION NUMBER: US 07/853,408
FILING DATE: 18-MAR-1992
PRIOR APPLICATION DATA:
                                                                                                                     APPLICATION NUMBER: US/07/853,408B FILING DATE: 19920318 CLASSIFICATION: 800
                                                                                                                                                                                  ATTORNEY AGENT INFORMATION:
NAME: Smith, William M.
REGISTRATION UNDRER: 30,223
REFERENCE/DOCKET 1464:
TELECOMMUNICATION INFORMATION:
TELEFONE: 415.326.2400
TELEFAX: 415.326.2420
INFORMATION FOR SEQ ID NO: 64:
SEQUENCE CHRARACTERISTICS:
LENGTH: 42 base pairs
TYPE: NUCLEIC ACID
STRANDENESS: 51091e
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 cctgacatcagtagcatctc 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      35 CCAGACATCAAAAGCATCAC 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear MOLECULE TYPE: DNA (primer)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
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TITLE OF INVENTION: Transgenic No. 6255458-Human Animals for TITLE OF INVENTION: Producing Heterologous Antibodies CORRESPONDENCE: 421
CORRESPONDENCE ADDRESS: ADDRESSE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco
                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/042,353
FILING DATE: 13-MAR-1998
CLASSIFICATION: 800
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 08/758,417
FILING DATE: 02-DEC-1996
FRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US97/21803
FILING DATE: 01-DEC-1997
ATORNEY/AGENT INFORMATION:
RAME: Apple, Randolph T.
REGISTRATION NUMBER: 36,429
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DAILS.
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US96/16433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/810,279
FILING DATE: 17-DEC-1991
PRIOR APPLICATION NUMBER: US 07/853,408
FILING DATE: 18-MAR-1992
PRIOR APPLICATION NATA:
APPLICATION NUMBER: US 07/900,860
FILING DATE: 23-0UN-1992
PRIOR APPLICATION NATA:
APPLICATION NUMBER: US 07/990,860
FILING DATE: 16-DEC-1992
PRIOR APPLICATION NATA:
APPLICATION NUMBER: US 08/053,131
FILING DATE: 22-0UL-1933
PRIOR APPLICATION NUMBER: US 08/155,301
FILING DATE: 18-NOV-1933
PRIOR APPLICATION NUMBER: US 08/155,301
FILING DATE: 03-DEC-1993
PRIOR APPLICATION NUMBER: US 08/165,699
FILING DATE: 10-DEC-1993
PRIOR APPLICATION NUMBER: US 08/165,699
FILING DATE: 10-DEC-1993
PRIOR APPLICATION NUMBER: US 08/165,699
FILING DATE: 10-DEC-1994
PRIOR APPLICATION DATA: APPLICATION NUMBER: US 08/352,322
FILING DATE: 07-DEC-1994
PRIOR APPLICATION DATA: APPLICATION NUMBER: US 08/352,322
FILING DATE: 07-DEC-1995
FILING DATE: 07-DEC-1995
PRIOR APPLICATION DATA: US 08/352,322
FILING DATE: 07-DEC-1995
PRIOR APPLICATION DATA: US 08/352,322
FILING DATE: 07-DEC-1995
PRIOR APPLICATION DATA: US 08/354,404
FILING DATE: 10-0CT-1995
PRIOR APPLICATION DATA: US 08/328,463
  Kay, Robert M
                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                 California
                                                                                                                                                                                                                                        COUNTRY: USA
ZIP: 94111-3834
APPLICANT:
                                                                                                                                                                                                                 STATE:
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APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transgenic No. 5877397-Human Animals for
TITLE OF INVENTION: Producing Heterologous Antibodies
CORRESPONDENCE: 150
CORRESPONDENCE ADDRESS:
                   Score 13.6; DB 2; Length 42;
Pred. No. 1.2e+02;
0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: William M. Smith
STREET: One Market Plaza, Steuart Tower, Suite 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: TIBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/308,865
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match

68.0%; Score 13.6; DB 2;
Best Local Similarity 80.0%; Pred. No. 1.2e+02;
Matches 16; Conservative 0; Mismatches 4;
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PRIOR APPLICATION DATA:
PPLICATION NUMBER: US/08/145,707
FILING DATE:
APPLICATION UNDHER: US 07/904,068
FILING DATE: 23-JUN-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.5-09-042-353-188/c

Sequence 188, Application US/09042353

Patent No. 6255458

SERERAL INFORMATION:

APPLICANT: Lonberg, Nils
                                                                                                                                                                                                                                                                                          US-08-308-865-64/c
; Sequence 64, Application US/08308865
; Betent No. 5877397
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Smith, William M.
REGISTRATION NUMBER: 30,223
REFERENCE/DOCKET NUMBER: 1464
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-326-2400
TELEFAX: 415-326-2422
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                                                                                                                             1 cctgacatcagtagcatctc 20
                                                                                                                                                                             35 CCAGACATCAAAAGCATCAC 16
                Query Match 68.0%;
Best Local Similarity 80.0%;
Matches 16; Conservative
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 42 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: One Market F
CITY: San Francisco
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
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US-08-308-865-64
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68.0%; Score 13.6; DB 4; 80.0%; Pred. No. 1.2e+02;
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APPLICATION UNBER: US 08/352,322
FILING DATE: 07-DEC-1994
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
FILING DATE: 10-OCT-1995
APPLICATION UNBER: US 08/728,463
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUBER: US 08/728,463
FILING DATE: 10-OCT-1996
PRIOR APPLICATION NUBER: WO PCT/US96/16433
FILING DATE: 10-OCT-1996
PRIOR APPLICATION NUBER: US 08/758,417
FILING DATE: 02-DEC-1996
PRIOR APPLICATION DATA:
APPLICATION NUBER: WO PCT/US97/21803
FILING DATE: 01-DEC-1997
ATPONEY AGAIN INFORMATION:
APPLICATION NUBER: WO PCT/US97/21803
FILING DATE: 01-DEC-1997
ATPONEY AGAIN INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Search completed: October 2, 2001, 16:03:50 Job time: 14594 sec
                                                              APPLICATION NUMBER: US 08/209,741
FILING DATE: 09-MAR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
INFORMATION FOR SEQ ID NO: 225:
SEQUENCE CHARACTERISTICS:
LENGTH: 42 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 cctgacatcagtagcatctc 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 68.0
Best Local Similarity 80.0
Matches 16; Conservative
                                                                                        FILING DATE: 09-MAR-PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; MOLECULE TYPE: DNA
US-09-042-353-225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        δ
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APPLICANT: Kay, Robert M.
TITLE OF INVENTION: Transgenic No. 6255458-Human Animals for
TITLE OF INVENTION: Producing Heterologous Antibodies
CORRESPONDENCE: 421
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                    Score 13.6; DB 4; Length 42;
Pred. No. 1.2e+02;
0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA

ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Bubbarcadero Center, Eighth Floor CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/09/042,353
FILING DATE: 13-MAR-1998
CLASSIPECATION: 800
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/810,279
FILING DATE: 17-DEC-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/853,408
FILING DATE: 18-MAR-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR DATE: 10-MAR 1992

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/904,068
FILING DATE: 23-UN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/990,860
FILING DATE: 16-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/053,131
FILING DATE: 26-APR-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/096,762
FILING DATE: 25-APR-1993
PRIOR APPLICATION UMBER: US 08/096,762
FILING DATE: 25-APR-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 08/096,762 FILING DATE: 22-UUL-1993 PRIOR APPLICATION DATA: APPLICATION NUMBER: US 08/155,301 FILING DATE: 18-NOV-1993
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03-DEC-1993
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; Sequence 225, Application US/09042353
; Detent No. 6255458
; CENERAL INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INPORMATION FOR SEQ ID NO: 188:
SEQUENCE CHARACTERISTICS:
LENGTH: 42 base pairs
TYPE: nucleic acid
STRANBEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 cctgacatcagtagcatctc 20
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                                                                                                                                                                                                                                                                                                                                             68.0%;
80.0%;
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Best Local Similarity 80.0
Matches 16; Conservative
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APPLICATION NUMBER: (
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                                                                                                                                                                                               ; TOPOLOGY: linear
; MOLECULE TYPE: DNA
US-09-042-353-188
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Length 42; Indels

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APPLICANT: Monital Brett P.
APPLICANT: Gaarde, William A.
APPLICANT: Garde, William A.
APPLICANT: Garde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0533
CURRENT APPLICATION NUMBER: US/09/757,100B
CURRENT APPLICATION NUMBER: 09/377,310
PRIOR FILING DATE: 1999-08-19
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STREET: 300 S. Wacker Drive Suite 3200
CITY: CHICAGO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/144,428
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 75.0%; Score 15; DB 29; 1
Best Local Similarity 100.0%; Pred. No. 7.2e+03;
Matches 15; Conservative 0; Mismatches 0;
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CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US97/03894
APPLICATION NUMBER: WO PCT/US97/03894
                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: PCT/US00/18999 PRIOR FLING DATE: 2000-07-13 NUMBER OF SEQ ID NOS: 44 SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: TAMBURINI, Paul P
APPLICANT: DAVIS, Gary
APPLICANT: DELARIA, Katherine A
APPLICANT: MARLOR, Christopher W
APPLICANT: MULLER, Daniel K
TITLE OF INVENTION: HUMAN BIKUNIN
NUMBER OF SEQUENCES: 71
CORRESPONDENCE ADDRESS:
ADDRESSEE: McDonnell Boehnen Hulbe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: antisense sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 31, Application US/09144428 GENERAL INFORMATION:
APPLICANT: BAYER CORPORATION, The
                                                                                                                                                                    ; Sequence 37, Application US/09757100B
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
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                                               1 tectegetgetggtg 15
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COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE:
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LENGTH: 15
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                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Isis Pharmaceuticals, Inc.
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
APPLICANT: Garde, William A.
APPLICANT: Nero, Pamela S.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase TITLE OF INVENTION: Expression
FILE REPERENCE: ISPH-0476
CURRENT APPLICATION NUMBER: PCT/USO0/18999
CURRENT FILING DATE: 2000-07-13
PRION FPLICATION NUMBER: 09/377,310
PRION FILING DATE: 1999-08-19
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APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
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; Pred. No. 7.2e+03;
0; Mismatches 0: TnAcl
                                                                                                                           Indels
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                                                                                76.0%; Score 15.2; DB 8;
85.0%; Pred. No. 6.1e+03;
Live 0; Mismatches 3;
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CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           75.0%; Score 15; 100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                               Sequence 37, Application PC/TUS0018999 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 37, Application US/09377310A GENERAL INFORMATION:
  DNA (genomic)
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Best Local Similarity 85.09
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Best Local Similarity 100.0
Matches 15; Conservative
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Best Local Similarity 100.
Matches 15; Conservative
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; MOLECULE TYPE:
US-08-466-588-52
                                                                                                                                                                                                                                                                          RESULT 5
PCT-US00-18999-37
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LENGTH: 15
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LENGTH: 15
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APPLICANT: Taylor, William J.A.

TITLE OF INVENTION: A method for Accelerating the Rate of Mucociliary Clearance FILE REFERENCE: 98,736-A
CURRENT APPLICATION NUMBER: US/09/441,966
CURRENT FILING DATE: 1999-11-17
PRIOR APPLICATION NUMBER: 09/218,913
PRIOR FILING DATE: 1998-12-22
NUMBER OF SEQ ID NOS: 71
SOFTWARE: Microsoft Word 97
SEQ ID NO 31
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GENERAL INFORMATION:
APPLICANT: Larry J. Smith
TITLE OF INVENTION: Methods and Compositions for Cellular
TITLE OF INVENTION: Reprogramming
FILE REFERENCE: Hesed-1
CURRENT APPLICATION NUMBER: US/08/668,235
CURRENT FILING DATE: 1996-06-17
EARLIER FILING DATE: 08/23/91
EARLIER FILING DATE: 08/23/91
EARLIER FILING DATE: 04/22/95
EARLIER FILING DATE: 04/22/95
EARLIER FILING DATE: 06/07/95
SEALIER FILING DATE: 06/07/95
NUMBER OF SEQ ID NOS: 3629
SOFTWARE: FastSEQ for Windows Version 3.0
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APPLICANT: Hesed 2.

APPLICANT: Smith, Larry J.

TITLE OF INVENTION: Method and Compositions for Cellular;

TITLE OF INVENTION: Reprogramming

TITLE OF INVENTION NUMBER: US/08/472,801

CURRENT FILING DATE: 1995-06-07

NUMBER OF SEQ ID NOS: 3601

SOFTWARE: FASTSEQ for Windows Version 3.0

SEQ ID NO 1314

LENGTH: 37
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Pred. No. 9.3e+03;
0; Mismatches 2;
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Pred. No. 1.4
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; GENERAL INFORMATION:
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93.8%;
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Best Local Similarity 88.9%;
Matches 16; Conservative
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Best Local Similarity 93.8
Matches 15; Conservative
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CORGANISM: S. cerevisiae
US-09-441-966-31
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US-08-472-801-1314
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GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Hall, Roderick L
APPLICANT: Newton, Benjamin B.
APPLICANT: Newton, Benjamin B.
TITLE OF INVENTION: A Method for Accelerating the Rate of Mucociliary Clearance FILE REFERENCE: 98,736
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Pred. No. 9.3e+03;
); Mismatches 2;
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Pred. No. 9.3e+03;
0; Mismatches 2;
                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/013,106
FILING DATE: 11-MAR-1996
PRIOR APPLICATION NUMBER: US 60/019,793
FILING DATE: US 60/019,793
FILING DATE: US 00/019,793
FILING DATE: US 08/725,251
FILING NAME: CHOO, MARK
NAME: CHAO, MARK
REFERENCE,DOCKET NUMBER: 96,223-II
TELECOMMUNICATION INFORMATION:
TELEPAX: (312) 913-0001
TELEPAX: (312) 913-0001
TELEFAX: (312) 913-0001
TELEPAX: (312) 913-0001
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CURRENT FILING DATE: 1998-12-22
NUMBER OF SEQ ID NOS: 71
SOFTWARE: Microsoft Word 97
SEQ ID NO 31
LENGTH: 35
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APPLICANT: Hall, Roderick L
APPLICANT: Poll, Christopher T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; TOPOLOGY: unknown
; MOLECULE TYPE: DNA (genomic)
US-09-144-428-31
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88.9%;
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88.9%;
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Best Local Similarity 88.9
Matches 16; Conservative
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Best Local Similarity 88.9
Matches 16; Conservative
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; ORGANISM: S. cerevisiae
US-09-218-9138-31
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US-09-218-913B-31
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US-09-441-966-31
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TYPE: DNA
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                                                                                                                                     Gaps
                                                                                                                                                            RESULT 13
PCT-US97-06104-16/C
; Sequence 16, Application PC/TUS9706104
; GENERAL INFORMATION:
; TITLE OF INVENTION: OLIGONUCLEOTIDES WITH ANTI-EPSTEIN-BARR
; TITLE OF INVENTION: VIGOS ACTIVITY
; TITLE OF INVENTION: VIGOS ACTIVITY
; NUMBER OF SEQUENCES: 43
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Fish & Richardson, P.C.
; STREET: 225 Franklin Street
; CITY: Boston
                                                                                                                                   ö
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                                                                                                Length 37;
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                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                              Score 14.4; DB 10;
Pred. No. 1.4e+04;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: ...
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM COMPATIBLE
OPERTING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US97/06104
FILING DATE: ...04 - APR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 14.2; DB 1;
Pred. No. 1.7e+04;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04020/094WO1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 132115, Application US/60232638 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08/628,422
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 05-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Fasse, J Peter
REGISTRATION NUMBER: 32,983
REFERENCE/DOCKET NUMBER: 0402/
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 617-542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        71.0%;
84.2%;
                                                                                                72.0%;
93.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ouery Match 71.0
Best Local Similarity 84.2
Matches 16; Conservative
                                                                                              Query Match 72.0
Best Local Similarity 93.8
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 04-APR-1
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: C
            TYPE: DNA
CORGANISM: Homo sapiens
US-08-668-235-1314
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; MOLECULE TYPE: DNA PCT-US97-06104-16
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US-60-232-638-132115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
LENGTH: 37
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US-60-234-017-326383/C
; Sequence 326383 Application US/60234017
; GENERAL INFORMATION:
; APPLICANT: Mittmann, M
; APPLICANT: Affymetrix, Inc.
; TITLE OF INVENTION: methods of Genetic Analysis of Mus
; TITLE OF INVENTION: methods of Senetic Analysis of Mus
; FILE REFERENCE: 3115
; CURRENT AFDLING DATE: 2000-09-20
; NUMBER OF SEQ ID NOS: 605887
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of Yeast FILE REFERENCE: 3110
CURRENT APPLICATION NUMBER: US/60/232,638
CURRENT FILING DATE: 2000-09-14
NUMBER OF SEQ ID NOS: 138410
SOFTWARE: FastSEQ for Windows Version 4.0
ENGTH: 25
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80.0%; Pred. No. 3.3e+04;
ive 0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                             68.0%; Score 13.6; DB 55
80.0%; Pred. No. 3.3e+04;
tive 0; Mismatches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Mus musculus
PUBLICATION INFORMATION:
DATABAGA ACCESSION NUMBER: Genbank AA153229
US-60-234-017-326383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 326383 LENGTH: 25
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                                                                                                                                                                                                                                                     ; PUBLICATION INFORMATION:
; DATABASE ACCESSION NUMBER: SGD YPL144W
US-60-232-638-132115
                                                                                                                                                                                                           TYPE: DNA ORGANISM: Saccharomyces Cerevisea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 attectegetgetggtggaa 20
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Best Local Similarity 80.0
Matches 16; Conservative
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Best Local Similarity 80.0
Matches 16; Conservative
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Job time: 24531 sec
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44, Appl 6820, Ap 2863, Ap 3510, Ap

Sequence Seq

App]

Sequence Seq

Searched:

Database

Result Š.

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Sequence:

Run on:

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APPLICANT: SANCHEZ, CESAR
APPLICANT: CHEN, MEI
APPLICANT: CHEN, MEI
TITLE NET BENERGE, DANIEL J.
TITLE OF INVENTOR: BLEOWICIN GENE CLUSTER COMPONENTS AND THEIR USES
FILE REFERENCE: 4077-895820US
CURRENT APPLICATION NUMBER: US/09/477,962
CURRENT APPLICATION NUMBER: 60/115,435
PRIOR FILING DATE: 1999-01-06
PRIOR FILING DATE: 1999-01-05
NUMBER OF SEQ ID NOS: 133
SOFTWARE: PATENTIN VEI. 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; OTHER INFORMATION: Description of Artificial Sequence: Primer US-09-477-962-58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       70.0%; Score 14; DB 6; Length 18; 100.0%; Pred. No. 1.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 39, Application US/09766450
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Collins, Collin
APPLICANT: Gray, Joe W.
APPLICANT: Albertson, Donna G.
APPLICANT: Pinkel, Daniel
APPLICANT: Pinkel, Daniel
APPLICANT: The Regents of the University of California
TITLE OF INVENTION: Repeat-Free Probes for Molecular
             US-09-708-690-4797

US-09-708-690-6943

US-09-708-690-6944

US-09-708-690-6945

US-09-708-690-7664

US-09-78-725-44

US-09-108-452A-6820

US-09-108-452A-6820

US-09-108-452A-4453

US-09-532-5378-5310

US-09-532-5378-54

US-09-531-700-111

US-09-801-16

US-09-801-174-122
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Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 58, Application US/09477962; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 70.0
Best Local Similarity 100.
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: SHEN, BEN
APPLICANT: DU, LIANGCHENG
APPLICANT: SANCHEZ, CESAR
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660.
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US-09-766-450-39/c
SEQ ID NO 58
LENGTH: 18
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Sequence 39, Appl
Sequence 5835, Ap
Sequence 27973, A
Sequence 6946, Ap
Sequence 6946, Ap
Sequence 6946, Ap
Sequence 5936, Ap
Sequence 6492, Ap
Sequence 6492, Ap
Sequence 6492, Ap
Sequence 6492, Ap
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638, App
4473, Ap
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3939, Ap
3115, Ap
160, App
5835, Ap
                                                                                                                              (without alignments)
19.290 Million cell updates/sec
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Sequence 5835, R
Sequence 4795, R
Sequence 4796, R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 6944,
Sequence 6944,
Sequence 6945,
Sequence 4795,
                                                                                                            ; Search time 2173.58 Seconds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Description
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Sequence 3
Sequence 3
Sequence 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pending_Patents_NA_New:*

1: /cgn2_6/ptodata/2/pna/PCT_NEW_COMB.seq:*

2: /cgn2_6/ptodata/2/pna/USO6_NEW_COMB.seq:*

3: /cgn2_6/ptodata/2/pna/USO8_NEW_COMB.seq:*

4: /cgn2_6/ptodata/2/pna/USO8_NEW_COMB.seq:*

5: /cgn2_6/ptodata/2/pna/USO9_NEW_COMB.seq:*

6: /cgn2_6/ptodata/2/pna/USO9_NEW_COMB.seq:*

7: /cgn2_6/ptodata/2/pna/USO9_NEW_COMB.seq:*

8: /cgn2_6/ptodata/2/pna/USO9_NEW_COMB.seq2:*
              GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-801-274-757
US-09-371-772B-6946
US-09-708-690-9123
US-09-708-690-9123
US-09-912-935-48
US-09-912-935-48
US-09-912-935-48
US-09-912-935-48
US-09-912-935-48
US-09-912-935-48
US-09-912-935-48
US-09-574-78-232-3939
US-09-574-78-5835
US-09-371-772B-4795
US-09-371-772B-4795
US-09-371-772B-4795
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US-09-371-772B-6944
US-09-371-772B-6945
US-09-708-690-4795
                                                                                                                                                                                                                                                                                                                  Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                    1895189 seqs, 1048201267 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SUMMARIES
                                                                                                            2001, 16:55:05
                                                                                                                                                                                                                                                                                                                                                                                              Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                            - nucleic search, using sw model
                                                                                                                                                                                                         1 attectegetgetggtggaa 20
                                                                                                                                                                                                                                       IDENTITY_NUC
Gapop 10.0 , Gapext 1.0
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20
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                                                                                                            October
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query
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112.8
112.8
112.6
112.4
12.4
12.4
12.5
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Gaps

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Indels

0 0 0 0 0 0 0 0 0

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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
APPLICANT: Pavco, Pam
APPLICANT: Stinochcomb, Dan
APPLICANT: Stinochcomb, Dan
APPLICANT: Stinochcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
CORRENT APPLICATION NUMBER: US/09/371,72B
CURRENT FILING DATE: 1995-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
PRIOR FILING DATE: 1996-01-08
NUMBER: OF SEQ ID NOS: 14225
SEQ ID NOS: 14225
SEQ ID NO 6946
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          Gaps
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                                                                                                                  RESULT 5
US-09-801-274-757/C
SQUENCE 757, Application US/09801274
SQUENCE 757, Application US/09801274
SQUENCE 757, Application US/09801274
SQUENCE 757, Application US/09801274
SPELICANT: Cargill, Michele
SAPPLICANT: Lander, Eric S.
TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
STILE REPRENCE: 2825.2009-001
CURRENT APPLICATION NUMBER: US/09/801,274
CURRENT FILING DATE: 2000-03-07
PRIOR FILING DATE: 2000-03-07
PRIOR FILING DATE: 2000-03-07
PRIOR FILING DATE: 2000-05-22
NUMBER OF SEQ ID NOS: 1802
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 757
TENGIN: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 17;
        Indels
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      1;
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Pred. No. 7e+03;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 13; DB 6;
Pred. No. 6e+03;
        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 6946, Application US/09371772B
; GENERAL INFORMATION:
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86.7%;
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Best Local Similarity 87.5%;
Matches 14; Conservative (
      Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 86.7
Matches 13; Conservative
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                                              3 tectegetgetggtg 17
|||| |||||||||||
22 tecttgetgetggtg 36
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; ORGANISM: Homo sapiens
US-09-371-772B-6946
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; ORGANISM: Homo sapiens
US-09-801-274-757
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US-09-371-772B-6946/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
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Sequence 5835, Application US/09198452A

GENERAL INFORMATION:

TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragments

TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, preverints of INVENTION: and treatment of infection

FILE REPERENCE: 9710-003-999

CURRENT APPLICATION NUMBER: US/09/198,452A

CURRENT FILING DATE: 1998-11-24

NUMBER OF SEQ ID NOS: 6849

SEQ ID NOS 6835
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GENERAL INFORMATION:
APPLICANT: Havukkala, Ilkka J
TILLE OF INVENTION: Polynucleotides, isolated from
TITLE OF INVENTION: Plants, and methods for their use.
FILE REFERENCE: 1054P1
CURRENT PILING DATE: 2000-11-27
NUMBER OF SEQ ID NOS: 37096
SEQ ID NOS: 37096
SEQ ID NO 27973
LENGTH: 44
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Pred. No. 3.6e+03;
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TITLE OF INVENTION: Cytogenetics FILE REFERENCE: 023071-111800US CURRENT APPLICATION NUMBER: US/09/766,450 CURRENT FILING DATE: 2001-01-19 NUMBER OF SEQ ID NOS: 112 SOFTWARE: FASTSEQ for Windows Version 3.0 SEQ ID NO 39
                                                                                                                                                                                                                                    ; OTHER INFORMATION: primer 727725.f1
US-09-766-450-39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
CRGANISM: Chlamydia pneumoniae
US-09-198-452A-5835
                                                                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence
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93.3%;
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93.3%;
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Best Local Similarity 93.3
Matches 14; Conservative
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Best Local Similarity 93.3
Matches 14; Conservative
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ORGANISM: Pinus radiata
US-60-253-456-27973
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Best Local Similarity
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RESULT 10
US-09-198-452A-6492
Sequence 6492, Application US/09198452A
GENERAL INFORMATION:
APPLICANT CALIffais, R.
TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, pr
TITLE OF INVENTION: and treatment of infection
TITLE OF INVENTION: and treatment of infection
CURRENT PILIAGO 1999
CURRENT APPLICATION NUMBER: US/09/198,452A
SEQ ID NOS: 6849
SEQ ID NO 6492
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                                                                      Length 17;
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                                                                                                                 Indels
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APPLICANT: Glenn, Murray R
TITLE OF INVENTION: Compositions isolated from bovine
TITLE OF INVENTION: Lissues and methods for their use.
FILE REPERENCE: 1052P2
CURRENT APPLICATION NUMBER: US/60/252,833
CURRENT FILING DATE: 2000-11-21
NUMBER OF SEQ ID NOS: 43535
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 5936
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 12.8; DB 8;
Pred. No. 7.8e+03;
0; Mismatches 2;
                                                                  Score 12.8; DB 5
Pred. No. 7e+03;
); Mismatches
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0S-60-252-833-5936/c
: Sequence 5936, Application US/60252833
; GENERAL INFORMATION:
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; ORGANISM: Chlamydia pneumoniae
US-09-198-452A-6492
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87.5%;
                                                                  64.0%;
87.5%;
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                                                               Query Match
Best Local Similarity 87.5
Matches 14; Conservative
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Best Local Similarity 78.9
Matches 15; Conservative
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Best Local Similarity
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; ORGANISM: Bovine
US-60-252-833-5936
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US-09-912-935-48/c
US-09-708-690-9123
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APPLICANT: Argued, Jam

APPLICANT: Stinchcomb, Dan

APPLICANT: Stinchcomb, Dan

APPLICANT: Stinchcomb, Dan

APPLICANT: Escobed, Jaiman Reagent for the Treatment of Diseases or Conditions Re

TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re

TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor

FILE REFERENCE: MBHB00,876-L (400/002)

CURRENT APPLICATION NUMBER: US 60/005,974

PRIOR APPLICATION NUMBER: US 60/005,974

PRIOR FILING DATE: 1996-01-08

PRIOR FILING DATE: 1996-01-08

PRIOR PILING DATE: 1999-08-10

PRIOR PILING DATE: 2000-10-10

PRIOR FILING DATE: 2000-10-10

NUMBER OF SEQ ID NOS: 20828

SOFTWARE: PatentIn Version 3.0

SEQ ID NO 6946
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
APPLICANT: Pavco, Pam
APPLICANT: Pavco, Pam
APPLICANT: Stanchcomb, Dan
APPLICANT: Stanchcomb, Dan
APPLICANT: Stanchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
CURRENT FILING DATE: 2001-08-31
PRIOR PLILING DATE: 1995-10-26
PRIOR PELICATION NUMBER: US 08/584,040
PRIOR PELICATION NUMBER: US 08/584,040
PRIOR PELING DATE: 1996-01-08
PRIOR FILING DATE: 1996-01-08
PRIOR FILING DATE: 1999-08-10
PRIOR PELING DATE: 2000-10-10
PRIOR PELING DATE: 2000-10-10
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Pred. No. 7e+03;
); Mismatches 2;
                                                                                                          Sequence 6946, Application US/09708690 GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-708-690-9123/C

3 Sequence 9123, Application US/09708690

3 GENERAL INFORMATION:
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SOFTWARE: PatentIn version 3.0
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87.5%;
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Best Local Similarity 87.5
Matches 14; Conservative
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17 CTTCCTGCTGGTGGAA
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                                                                                  US-09-708-690-6946/c
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LENGTH: 17
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OTHER INFORMATION: Description of Artificial Sequence: Nucleic Acid
NAME/KEY: misc_feature
LOCATION: (17)..(26)
OTHER INFORMATION: positions 17-26 n stands for any nucleotide, positions 19-26 n
OTHER INFORMATION: be optionally absent
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Stinchcomb, Dan
APPLICANT: Draper, Ken
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Rel
TITLE OF INVENTION: Levels of NF-Kappa B
FILLE REFERENCE: 400/022 (MBHB00-812-D)
CURRENT APPLICATION NUMBER: US/09/864,785
CURRENT FILING DATE: 2001-05-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 3939, Application US/60278232
GENERAL INFORMATION:
APPLICANT: Morris, MacDonald
APPLICANT: Lal, Preeti
APPLICANT: Disp, Dinh
TITLE OF INVENTION: Polymorphisms Identified Thereby
TITLE OF INVENTION: Polymorphisms Identified Thereby
CURRENT APPLICANTON NUMBER: US/60/278,232
CURRENT FILING DATE: 2001-03-30
NUMBER OF SEQ ID NOS: 12,557
SOFTWARE: PERL PROGram
SEQ ID NO 3939
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Pred. No. 1.2e+04;
0; Mismatches 1;
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US-60-278-232-3939
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; GENERAL INFORMATION:
         SOFTWARE: PatentIn version 3.0 SEQ ID NO 4473 LENGTH: 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            62.0%;
92.9%;
                                                                            TYPE: RNA
ORGANISM: Artificial Sequence
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64.3%;
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Best Local Similarity 64.3
Matches 9; Conservative
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ORGANISM: Homo sapiens
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Best Local Similarity
Matches 13; Conserv
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APPLICANT: Nishikawa, Mitsuo et al.

TITLE OF INVENTION: METHODS AND MATERIALS RELATING TO STEM CELL GROWTH FACTOR-LIKE
TITLE OF INVENTION: POLYPEPTIDES AND POLYNUCLEOTIDES
FILE REPERENCE: 32066/37483
CURRENT PILING DATE: 2001-07-24
PRIOR PILING DATE: 2001-12-23
NUMBER OF SEQ ID NOS: 53
SOFTWARE: PALENTIN VERSION 3.0
ESQ ID NO 48
LENGTH: 24
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APPLICANT: Blatt, Larry
APPLICANT: Blatt, Larry
APPLICANT: Ewick, Michael
APPLICANT: Pavco, Pam
APPLICANT: McSatiggen, Jim
TITLE OF INVENTION: Regulation of Repressor Genes using Nucleic Acid Molecules
FILE REFERENCE: 237/193
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GENERAL INFORMATION:
APPLICANT: Cargill, Michele
APPLICANT: Lander, Eric S.
TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
FILE REFERENCE: 2825.2009-001
CURRENT APPLICATION NUMBER: US/09/801,274
CURRENT FILING DATE: 2001-03-07
PRIOR APPLICATION NUMBER: US 60/187,510
PRIOR FILING DATE: 2000-03-07
PRIOR FILING DATE: 2000-05-22
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Pred. No. 1.2e+04;
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Pred. No. 9.1e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 1802
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 638
LENGTH: 31
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78.9%;
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81.2%;
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Best Local Similarity 78.9
Matches 15; Conservative
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Best Local Similarity 81.2
Matches 13; Conservative
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; ORGANISM: primer H2728F
US-09-912-935-48
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ORGANISM: Homo sapiens
US-09-801-274-638
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; NUMBER OF SEQ ID NOS: 3929
; SOFTWARE: PatentIn version 3.0
; SOFTWARE: PatentIn version 3.0
; SED ID NO 3115
; LENGTH: 48
; TYPE: RNA
; ORGANISM: Artificial Sequence
; FRATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Nucleic Acid
US-09-864-785-3115

Query Match
G2.0%; Score 12.4; DB 7; Length 48;
Best Local Similarity 78.6%; Pred. No. 1.2e+04;
Matches 11; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 7 cgctgctggtggaa 20
| | | | | | | | | | | | | |
| Db 2 cgcugcuggaggaa 15

Search completed: October 2, 2001, 16:55:05
Job time: 17668 sec
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RESULT 2
US-09-922-146-6/c
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18238, A
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Sequence 6, Appli
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                                                                                                                                                         Search time 2173.58 Seconds
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1: /cgn2_6/ptodata/2/pna/PCT_NEW_COMB.seq:*

2: /cgn2_6/ptodata/2/pna/USO6_NEW_COMB.seq:*

3: /cgn2_6/ptodata/2/pna/USO8_NEW_COMB.seq:*

4: /cgn2_6/ptodata/2/pna/USO8_NEW_COMB.seq:*

5: /cgn2_6/ptodata/2/pna/USO9_NEW_COMB.seq:*

6: /cgn2_6/ptodata/2/pna/USO9_NEW_COMB.seq:*

7: /cgn2_6/ptodata/2/pna/USO9_NEW_COMB.seq:*

8: /cgn2_6/ptodata/2/pna/USO9_NEW_COMB.seq2:*

8: /cgn2_6/ptodata/2/pna/USO9_NEW_COMB.seq2:*
                                           Compugen Ltd
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US-09-922-146-6
US-09-814-577-8
US-09-814-879-5909
US-09-708-690-18238
US-09-813-152-6
US-09-574-376B-1160
US-09-574-376B-1160
US-09-574-376B-1160
US-09-574-376B-1160
US-09-574-376B-1160
US-09-574-376B-1160
US-09-574-376B-1160
US-09-514-67-326
US-09-514-67-326
US-09-310-735A-140
US-09-310-735A-140
US-09-310-735A-151
US-09-310-844B-140
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US-09-421-971-19
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US-09-941-992-470
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                   GenCore version 4.5
Copyright (c) 1993 - 2000 Comp
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                                                                                                                                                         2001, 16:55:04
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Listing first 45 summaries
                                                                                                             OM nucleic - nucleic search, using sw model
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ALIGNMENTS

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RESULT 1

Sequence 1052, April 1052/C

1959-650-012A-1062/C

1959-650-012A-1062/C

Sequence 1052, April 1051.

Sequence 1052, April 1051.

Sequence 1052, April 1051.

APPLICANT: Riboyem, 1051.

APPLICANT: Strinchcomb, Danie Pharmaceuticals, Inc.

APPLICANT: Strinchcomb, Danie Pharmaceuticals, Inc.

APPLICANT: Strinch The Probability Thate

APPLICANT: Marker Pharmaceuticals, Inc.

STRING PARE: 1090-00-28

FRIOR PLING APPLICATION NUMBER: US 60/200.

FRIOR APPLICATION NUMBER: US 60/200.

SEQ 1D NO. 1062

SEQ 1D NO. 1063

SEQ 1064

SEQ
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APPLICANT: Study gen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stobedo, Jaime
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
CURRENT APPLICATION NUMBER: US/09/708, 590
CURRENT APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 09/371,772
PRIOR APPLICATION NUMBER: US 09/371,772
PRIOR APPLICATION NUMBER: US 09/371,772
PRIOR PRILING DATE: 1999-08-10
NUMBER OF SEQ ID NOS: 20828
SOFTWARE: PARENTIN VARIATION NUMBER: US 09/685,664
PRIOR FILING DATE: 2000-10-10
NUMBER OF SEQ ID NOS: 20828
SEQ ID NO 18238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: Description of Artificial Sequence: Enzymatic Nucleic Acid
US-09-708-690-18238
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Sequence 6, Application US/09942325
Sequence 6, Application US/09942325
GENERAL INFORMATION:
APPLICANT: Iacovitti, Lorraine
APPLICANT: Ressler, Mark
TITLE OF INVENTION: The Human Tyrosine Hydroxlyase Promoter
TITLE OF INVENTION: Sequence and Related Methods and Compositions
FILE REFERENCE: IACO1.NB01
CURRENT APPLICATION NUMBER: US/09/942,325
CURRENT PEPLICATION NUMBER: 60/228931
PRIOR FILING DATE: 2000-02-30
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                                                                                                                         Length 31;
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; 2.2e+04;
5; Indels
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Pred. No. 1.7e+04;
0; Mismatches 3;
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 6
LENGTH: 38
                                                                                                                                                                                                                                                                                                                                                                                ; Sequence 18238, Application US/09708690
; GENERAL INFORMATION:
; APPLICANT: Ribozyme Pharmaceuticals, Inc.;
APPLICANT: Pavco, Pam
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82.4%;
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                     ; OTHER INFORMATION: DNAZyme
US-09-817-879-5909
                                                                                                                  Query Match 61.0°
Best Local Similarity 82.4°
Matches 14; Conservative
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Best Local Similarity 60.0°
Matches 12; Conservative
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    LOCATION:
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GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals Inc.
TITLE OF INVENTION: Enzymatic Nucleic Acid Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Hepatitis C Virus Infection
FILE REFERENCE: MBHB00-801-F
CURRENT APPLICATION NUMBER: US/09/817,879
CURRENT APPLICATION NUMBER: US/09/817,879
SOFTWARE: 2001-26
NUMBER OF SEQ ID NOS: 9703
SOFTWARE: Patentin version 3.0
SEQ ID NO 5909
LENGTH: 31
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APPLICANT: Zhou, Hua
APPLICANT: Zhou, Hua
APPLICANT: Zhou, Hua
APPLICANT: Greene L., Amy
APPLICANT: Thode, Silke
TITLE OF INVENTION: METHODS AND NUCLEIC ACID VECTORS FOR RAPID AND PARALLEL
TITLE OF INVENTION: ASSAY DEVELOPMENT, FOR CHARACTERIZATION OF THE
TITLE OF INVENTION: ASSAY DEVELOPMENT, FOR CHARACTERIZATION OF THE
FILE REFERENCE: 5050-0015
CURRENT APPLICATION NUMBER: 05/99/545,777
CURRENT FILING DATE: 2000-04-07
PRIOR FILING DATE: 1999-04-09
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US-09-545-777-8
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                                                                                                                                                                                                                    Length 26;
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                                                                                                                                                                                                                                                                  4; Indels
                                                                                                                                                                                                                 Score 12.6; DB 7;
Pred. No. 1.1e+04;
                                                                                                                                                                                                                                                                0; Mismatches
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                                         TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                            Query Match 63.0%;
Best Local Similarity 78.9%;
Matches 15; Conservative (
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ORGANISM: artificial sequence
FEATURE:
NAME/KEY: misc_feature
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                                                                                          ; FEATURE:
; OTHER INFORMATION: PCR Probe
US-09-922-146-6
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SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 8
LENGTH: 32
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Best Local Similarity 78.9
Matches 15; Conservative
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SEQ ID NO 6
LENGTH: 26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            59.0%; Score 11.8; DB 5; Length 31; 76.5%; Pred. No. 2.7e+04;
Live 1; Mismatches 3; Indels
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                                                                                                                                                  Sequence 1320, Application US/09574376B
GENERAL INFORMATION:
APPLICANT: Warrington, Janet
APPLICANT: Gladen, Vila
APPLICANT: Gladeras, Thomas Raymond
APPLICANT: Affymetrix, Inc.;
TITLE OF INVENTION: Human Lymphoblast Polymorphisms
FILE REFERENCE: 3229, 2
CURRENT APPLICATION NUMBER: US/09/574,376B
CURRENT FILIG DATE: 2000-05-19
NUMBER OF SEQ ID NOS: 1330
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1320
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86.7%; Pred. No. 2.7e+04;
11ve 0; Mismatches 2;
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CURRENT FILING DATE: 2001-01-27
PRIOR APPLICATION NUMBER: 60/179,006
PRIOR FILING DATE: 2000-01-28
NUMBER OF SEQ ID NOS: 30
SOFTWARE: Patentin Ver: 2.1
SEQ ID NO 9
LENGTH: 39
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APPLICANT: POLLART, DAN
APPLICANT: SHALER, THOMAS
APPLICANT: STEPHENS, KATHY
APPLICANT: TAN, YUPING
APPLICANT: WONG, LINDA
APPLICANT: WONG, LINDA
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ORGANISM: Artificial Sequence
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                      1 ctaggggaggctcagtg 17
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14 ctmggggtcggtcagtg 30
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Matches 13; Conservative
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US-09-574-376B-1320
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Best Local Similarity
Matches 13; Conserv
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US-09-574-376B-1320
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APPLICANT: Reinhard, Christoph
APPLICANT: Reinhard, Christoph
APPLICANT: Jefferson, Anne Bennett
TITLE OF INVENTION: POLYNUCLEOTIDES RELATED TO COLON CANCER
FILE REFERENCE: 2300-1663
CURRENT FAPLICATION NUMBER: US/09/883,152
CURRENT FILING DATE: 2001-06-15
PRIOR RIDNG DATE: 2000-06-15
NUMBER OF SEQ ID NOS: 127
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 59
LENGTH: 22
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Pred. No. 2.7e+04;
1; Mismatches 3; Indels
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                                                                                                                                  0; Indels
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GENERAL INFORMATION:
APPLICANT: Warrington, Janet
APPLICANT: Shah, Nila
APPLICANT: Gingeras, Thomas Raymond
APPLICANT: Affymetrix, Inc.
TILLE OF INVENTION: Human Lymphoblast Polymorphisms
FILE REFERENCE: 3229.2
CURRENT APPLICATION NUMBER: US/09/574,376B
CURRENT FILING DATE: 2000-05-19
NUMBER OF SEQ ID NOS: 1330
SOFTWARE: FastSEQ for Windows Version 4.0
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Pred. No. 2.7e+04;
0; Mismatches 2;
                                                                                        Score 12; DB 5; L
; Pred. No. 2.2e+04;
0; Mismatches 0:
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Sequence 59, Application US/09883152
GENERAL INFORMATION:
APPLICANT: Kennedy, Giulia
                                                                                            60.0%;
100.0%;
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76.5%;
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86.7%;
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Best Local Similarity 76.55
Matches 13; Conservative
                                                                                            Query Match 60.0
Best Local Similarity 100.
Matches 12; Conservative
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Best Local Similarity 86.7
Matches 13; Conservative
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PUBLICATION INFORMATION:
US-09-574-376B-1160
; TYPE: DNA
; ORGANISM: Homo sapien
US-09-942-325-6
                                                                                                                                                                    US-09-574-376B-1160
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LENGTH: 31
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Application US/09514673
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Best Local Similarity 77.8
Matches 14; Conservative
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Matches 14; Conservative
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US-09-310-735A-140/c
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US-09-563-794-69/c
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US-09-786-033-19
Sequence 19, Application US/09786033
GENERAL INFORMATION:
APPLICANT: Pausch, Mark Henry
APPLICANT: Pausch, Mark Henry
TILLE OF INVENTION: METHODS FOR IMPROVING THE FUNCTION OF HETERLOGOUS G
TILLE OF INVENTION: METHODS FOR IMPROVING THE FUNCTION OF HETERLOGOUS G
TILLE OF INVENTION: METHODS FOR IMPROVING THE FUNCTION OF HETERLOGOUS G
TILLE OF INVENTION: METHODS FOR IMPROVING THE FUNCTION OF HETERLOGOUS G
TURRENT APPLICATION UNMBER: US/09/786,033
CURRENT FILING DATE: 2001-07-03
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 19
LEMOTH: 31
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                                                                                                       APPLICANT: Anthony, James
APPLICANT: Anthony, James
APPLICANT: Anthony, James
APPLICANT: Milliams, Inna
APPLICANT: Torio, Yanlin
APPLICANT: Trong, Yanlin
TITLE OF INVENTION: DETECTION OF NUCLEIC ACIDS BY TYPE-SPECIFIC HYBRID
TITLE OF INVENTION: DETECTION OF NUCLEIC ACIDS BY TYPE-SPECIFIC HYBRID
TITLE OF INVENTION: DETECTION OF NUCLEIC ACIDS BY TYPE-SPECIFIC HYBRID
TITLE OF INVENTION: DETECTION OF NUMBER: POT/USO1/19353
CURRENT APPLICATION NUMBER: POT/USO1/19353
CURRENT FILING DATE: 2000-06-15
NUMBER OF SED ID NOS: 162
SOFTWARE: Patentin Ver. 2.1
SED ID NO 77
LENGTH: 31
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OTHER INFORMATION: Description of Artificial Sequence:nucleic acid
CTHER INFORMATION: probe
PCT-US01-19353-77
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Pred. No. 3.4e+04;
0; Mismatches 4; Indels
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                                                                 Sequence 77, Application PC/TUS0119353 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Artificial Sequence
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77.8%;
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ORGANISM: Artificial Sequence
FEATURE:
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Best Local Similarity 77.8
Matches 14; Conservative
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Matches 14; Conservative
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US-09-514-673-26/c
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TITLE OF INVENTION: NOVEL TRANSDUCTION MOLECULES AND METHODS FOR USING SAME FILE REPERENCE: 49054 (17142) CURRENT APPLICANTION NUMBER: US/09/514,673 CURRENT FILING DATE: 2000-02-28 PRIOR PILING DATE: 2000-02-28 PRIOR PILING DATE: 1999-02-28 PRIOR FILING DATE: 1999-02-29 PRIOR FILING DATE: 1999-02-29 RIOR PILING DATE: 1999-02-29 PRIOR FILING DATE: 1999-02-29 PRIOR FILING DATE: 1999-02-29 PRIOR PILING DATE: 1999-02-29 PRIOR PILING DATE: 1999-08-29 SOFTWARE: PATENTING DATE: 1990-08-29 SOFTWARE: PATENTING DATE: 1990-08-29 SOFTWARE: PATENTING DATE: 2.1
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GENERAL INFORMATION:
APPLICANT: Kruger, Martin
APPLICANT: Welch, Peter J.
APPLICANT: Welch, Peter J.
TITLE OF INVENTION: Cellular Regulators of Infectious Agents and Methods of TITLE OF INVENTION: Use
FILE REFERENCE: P-IU 3738
CURRENT APPLICATION NUMBER: US/09/563,794
CURRENT FILING DATE: 2000-05-02
NUMBER OF SEQ ID NOS: 140
SSOTWARE: PatentIn Ver. 2.1
SSOTUB NO 69
LIENGTH: 40
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Pred. No. 3.4e+04;
0; Mismatches 4; Indels
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GENERAL INFORMATION:
APPLICANT: Ecker, David J
APPLICANT: Griffey, Richard
APPLICANT: Grobe, Stanley T
APPLICANT: Sampath, Ranga
APPLICANT: Swayze, Eric
```

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## APPLICANT: Mohan, Venkatraman
## APPLICANT: Hofstadler, Steven
## APPLICANT: Hofstadler, Steven
### APPLICANT: Hofstadler, Steven
### APPLICANT: Hofstadler, Steven
### TITLE OF INVENTION: Other Biomolecules
### TITLE OF INVENTION: Other Biomolecules
### CURRENT PERLICANT: USP9-05-12
### CURRENT PELLOR DATE: 1999-05-12
### PRIOR APPLICATION OF APPLICATION
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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15 29 US-09-757-100B-36 Sequence 7185 4.4 72.0 25 26 US-09-660-220-7185 Sequence 7185 4.4 72.0 25 26 US-09-660-220-7185 Sequence 7185 3.8 69.0 25 55 US-60-233-166-38586 Sequence 1209 3.8 69.0 25 55 US-60-234-017-12091 Sequence 1209 3.8 69.0 25 55 US-60-234-017-12091 Sequence 1209 3.8 69.0 25 55 US-60-240-94301 Sequence 1209 3.8 69.0 25 55 US-60-224-017-12091 Sequence 1209 3.8 69.0 25 55 US-60-261-18 Sequence 1209 3.4 67.0 34 7 US-08-355-684-2592 Sequence 57, Mg 3.2 66.0 18 9 US-08-355-684-2592 Sequence 57, Mg 3.2 66.0 21 US-08-356-684-2592 Sequence 57, Mg 3.2 66.0 21 <		15	5	1		-09-377-310-36	Sequence 36,
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3.8 69.0 25 50 US-60-223-166-385865 Sequence 4064(35.8) 8.9 69.0 25 55 US-60-233-166-386865 Sequence 120933.8 69.0 25 55 US-60-234-017-120917 Sequence 120933.8 69.0 25 55 US-60-234-017-120917 Sequence 120933.8 69.0 25 55 US-60-234-017-120917 Sequence 120933.8 69.0 25 55 US-60-220-94301 Sequence 120933.4 67.0 34 7 US-08-305-7714-57 Sequence 57, April 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0 1.0		₹ <	٠. د	7 (-09-660-220-7185	equence 7185
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3.8 69.0 25 55 US-60-214-017-120921 Sequence 1209; 3.8 69.0 28.1 PCT-USOO-26619-18 Sequence 18.7 Mg		13.8	6	7		-60-234-017-12091	equence 1209
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3.4 66.0 18 9 US-08-359-7/14-5/7 Sequence 2592, 3.2 66.0 18 9 US-08-555-684-2592 Sequence 2592, 3.2 66.0 18 9 US-08-555-684-2592 Sequence 2592, 3.2 66.0 21 IP US-09-266-682-6 Sequence 6, Agrace 6.0 21 I7 US-09-322-134-6 Sequence 6, Agrace 6.0 21 I7 US-09-322-134-6 Sequence 6, Agrace 6.0 23 II US-08-422-134-6 Sequence 3446, 3.2 66.0 23 II US-08-68-233-166-23415 Sequence 23415 Sequence 275, Agrace 6.0 25 55 US-60-233-620-3638 Sequence 275, Agrace 6.0 33 II PCT-US96-0388-7 Sequence 275, Agrace 6.0 33 II PCT-US96-034-7 Sequence 275, Agrace 6.0 47 II US-09-245-024-7 Sequence 275, Agrace 6.0 47 II US-09-245-034-7 Sequence 275, Agrace 6.0 47 II US-09-245-034-7 Sequence 275, Agrace 6.0 47 II US-09-245-034-7 Sequence 275, Agrace 6.0 47 II US-09-245-034-13 Sequence 275, Agrace 6.0 47 II US-09-245-034-13 Sequence 275, Agrace 6.0 47 II US-09-245-034-13 Sequence 275, Agrace 6.0 49 II US-09-09-4904-94 Sequence 275, Agrace 6.0 49 II US-09-660-220-5688 Sequence 240, Agrace 6.0 25 55 US-60-220-10044 Sequence 240, Agrace 6.0 25 55 US-00-09865-13 Sequence 240, Agrace 6.0 25 55 US-00-233-166-24015II Sequence 10816 6.0 25 US-00-233-166-24015II Sequence 10816 6.0 25 US-00-233-166-24015II		13.4	٠,	34	~ 1	305-771A-5	53, Ap
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3.2 66.0 21 17 US-09-222-134-6 Sequence 3.446, 3.2 66.0 23 10 US-08-422-1345 Sequence 3.446, 3.2 66.0 23 10 US-08-68-235-3446 Sequence 3.446, 3.2 66.0 25 55 US-60-233-166-23415 Sequence 2.3411 Sequence 2.3411 Sequence 2.3411 Sequence 2.3411 Sequence 2.3412 Sequence 2.34		ກຕ	• u	2 50		9-23203-8 3-283-285	, , s
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		12.8	64.0	25		-60-234-017-3094	equence 3094

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Query Match
Best Local Similarity
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PCT-US00-18999-36
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LENGTH: 15
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                                                                                                                                               APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
APPLICANT: Nero, Pamela S.
TITLE OF INVENTION: ALISENSE Modulation of Focal Adhesion Kinase TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0476
CURRENT APPLICATION NUMBER: PCT/US00/18999
CURRENT FILING DATE: 2000-07-13
PRIOR APPLICATION NUMBER: 09/377,310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310A
CURRENT APPLICATION NUMBER: US/09/377,310A
SUMBER OF SEQ ID NOS: 43
SEQ ID NO 16
LENTH: 20
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Pred. No. 14;
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ALIGNMENTS
                                                                                                                                    Isis Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTHER INFORMATION: antisense sequence PCT-US00-18999-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CTHER INFORMATION: antisense sequence US-09-377-310-16
                                                                                              Sequence 16, Application PC/TUS0018999 GENERAL INFORMATION:
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US-09-757-100B-16
; Sequence 16, Application US/09757100B
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                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 16
LENGTH: 20
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Best Local Similarity 100.
Matches 20; Conservative
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Matches 20; Conservative
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US-09-377-310-16
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APPLICANT:
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APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Monia, Milliam A.
APPLICANT: Monia, Pamela S.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-6533
CURRENT APPLICATION NUMBER: US/09/757,100B
CURRENT FILING DATE: 12099-08-19
PRIOR APPLICATION NUMBER: 09/377,310
PRIOR FILING DATE: 12099-08-19
PRIOR FILING DATE: 2000-07-13
NUMBER OF SEQ ID NOS: 44
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 36, Application PC/TUS0018999
GENERAL INFORMATION:
APPLICANT: Isis Pharmaceuticals, Inc.
APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Morey, Pamela S.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase;
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0476
CURRENT APPLICATION NUMBER: PCT/US00/18999
CURRENT FILING DATE: 2000-07-13
PRIOR PILING DATE: 1999-08-19
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ) OTHER INFORMATION: antisense sequence PCT-US00-18999-36
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APPLICANT: Monia, Brett P.
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100.0%;
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SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20; Conservative
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Best Local Similarity 100.
Matches 15; Conservative
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APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of the Rat
TITLE OF INVENTION: Genome
FILE REFERENCE: 3112
CURRENT APPLICATION UNBER: US/60/233,166
CURRENT FILING DATE: 2000-10-24
NUMBER OF SEQ ID NOS: 420907
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APPLICANT: Affymerrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of
FILE REPERENCE: 3102.1
CURRENT APPLICATION NUMBER: US/09/660,220
CURRENT FILING DATE: 2000.09-12
PRIOR PLICATION NUMBER: 60/164,973
PRIOR FILING DATE: 1999.11-11
                                                                                                                                                                                                                                                                      Score 14.4; DB 26;
Pred. No. 7.6e+03;
0; Mismatches 1;
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93.8%; Pred. No. 7.6e+03;
live 0; Mismatches 1;
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US-60-233-166-385862
                             SEQ ID NOS: 140981
FastSEQ for Windows Version 4.0
                                                                                                              TYPE: DNA
ORGANISM: Homo Sapiens
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank S76992
US-09-660-220-71852
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SOCYTHARE: FastSEQ for Windows Version 4.0
SEQ ID NO 78149
LENGTH: 25
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PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank U10439
US-09-660-220-78149
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; Sequence 385862, Application US/60233166
; GENERAL INFORMATION:
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         1999-11-11
                                                                                                                                                                                                                                                                        72.0%;
93.8%;
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Best Local Similarity 93.8
Matches 15; Conservative
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Best Local Similarity
Matches 15; Conserv
         PRIOR FILING DATE:
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US-09-660-220-78149/c
                             NUMBER OF SEQ ID N
SOFTWARE: FastSEQ
SEQ ID NO 71852
LENGTH: 25
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GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
APPLICANT: Garade, William A.
TITLE OF INVENTION: Expression
FILE REFERENCE: ESPH-0533
CURRENT APPLICATION NUMBER: US/09/757,100B
CURRENT PILING DATE: 2001-03-15
PRIOR APPLICATION NUMBER: 09/377,310
PRIOR APPLICATION NUMBER: PCT/US00/18999
PRIOR PILING DATE: 1999-08-19
PRIOR APPLICATION NUMBER: PCT/US00/18999
PRIOR APPLICATION NUMBER: PCT/US00/18999
SROUP DATE: 2000-07-13
NUMBER OF SEQ ID NOS: 44
SSEQ ID NO 36
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310A
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver: 2.0
SEQ ID NO 36
LENGTH: 15
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GENERAL INFORMATION:
APPLICANT: Mittmann et al.
APPLICANT: Affmetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of Human FILE REFERENCE: 3102.1
CURRENT PAPLICATION NUMBER: US/09/660,220
CURRENT FILING DATE: 2000-09-12
PRIOR APPLICATION NUMBER: 60/164,973
                                                                                                                                                                                                                                                                                                                                                                               75.0%; Score 15; DB 100.0%; Pred. No. 3.7 tive 0; Mismatches
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Best Local Similarity 100.
Matches 15; Conservative
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Best Local Similarity 100.
Matches 15; Conservative
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-09-757-100B-36
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Sequence 18, Application PC/TUS0026619
; Sequence 18, Application PC/TUS0026619
; GENERAL INFORMATION:
; APPLICANT: Chmura, Albert
; APPLICANT: Chmura, Albert
; APPLICANT: Chmura, Albert
; TITLE OF INVENTION: Engineering Antibodies That Bind Irreversibly
; TITLE OF INVENTION: 203070-099120PC
; CURRENT PAPLICATION NUMBER: PCT/US00/26619
; CURRENT FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: US 60/156,194
; PRIOR APPLICATION NUMBER: US 60/208,684
; PRIOR APPLICATION NUMBER: US 60/208,684
; PRIOR APPLICATION NUMBER: US 60/208,684
; PRIOR PETLING DATE: 2000-05-31
; SOFTWARE: PatentIN Ver. 2.1
; SEQ. ID NO 18
; LENGTH: 28
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 25;
               Sequence 120921, Application US/60234017
GENERAL INFORMATION:
APPLICANT: Mittmann, M
APPLICANT: Afforetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of Mus
TITLE OF INVENTION: musculus
FILE REFERENCE: 3115
CURRENT APPLICATION NUMBER: US/60/234,017
CURRENT FILING DATE: 2000-09-20
NUMBER OF SEQ ID NOS: 605887
SEQ ID NO 120921
LENGTH: 25
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Pred. No. 1.5e+04;
0; Mismatches 2;
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Pred. No. 1.5e+04;
0; Mismatches 2;
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; DATABASE ACCESSION NUMBER: GenBank U64445
US-60-234-017-120921
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milarity 88.2%;
Conservative 0
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88.2%;
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Best Local Similarity 88.2
Matches 15, Conservative
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Best Local Similarity
Matches 15; Conserv
US-60-234-017-120921
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                 Length 25;
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                                                                                                                                                                                                                                                 Sequence 406405, Application US/60233166
GENERAL INFORMATION:
APPLICANT: Mittmann
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of the Rat
TITLE OF INVENTION: Genome
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APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of Mus.
TITLE OF INVENTION: musculus
FILE REPERENCE: 3115
CURRENT APPLICATION NUMBER: US/60/234,017
CURRENT FILING DATE: 2000-09-20
NUMBER OF SEQ ID NOS: 605887
SEQ ID NO: 605887
SEQ ID NO: 120917
LENGTH: 25
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88.2%; Pred. No. 1.5e+04;
tive 0; Mismatches 2;
                 Score 13.8; DB 55;
Pred. No. 1.5e+04;
0; Mismatches 2;
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Pred. No. 1.5e+04;
0; Mismatches 2;
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PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Affymetrix Proprieta
US-60-233-166-406405
                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: 3112
CURRENT APPLICATION NUMBER: US/60/233,166
CURRENT FILING DATE: 2000-10-24
NUMBER OF SEQ ID NOS: 420907
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 406405
LENGTH: 25
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; DATABASE ACCESSION NUMBER: GenBank U64445
US-60-234-017-120917
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US-60-234-017-120917.
; Sequence 120917, Application US/60234017
; GENERAL INFORMATION:
               Query Match 69.0%;
Best Local Similarity 88.2%;
Matches 15; Conservative
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88.2%;
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Best Local Similarity
Matches 15; Conserv
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APPLICANT: Mittmann et al. APPLICANT: Affymetrix, Inc.

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Length 34;
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Pred. No. 2.4e+04;
0; Mismatches 1;
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                                                    TYPE: nucleic acid
syrANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
US-08-305-771A-53
                                                                                                                                                                                                                         67.0%;
93.3%;
                           1: 34 base pairs
nucleic acid
DEDNESS: single
       SEQUENCE CHARACTERISTICS:
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Best Local Similarity
Matches 14; Conserva
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APPLICANT: Lois Chandler
APPLICANT: Michael Nova
TITLE OF INVENTION: GROWTH FACTOR-LIKE GROWTH FACTOR AND TARGETED AGENTS
NUMBER OF SEQUENCES: 87
CORRESPONDENCE ADDRESS:
ADDRESSEE: Brown, Martin, Haller & McClain
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TITLE OF INVENTION: Methods of Genetic Analysis of Human FILE REFERENCE: 3102.1
CURRENT APPLICATION NUMBER: US/09/660,220
CURRENT FILING DATE: 2000-09-12
PRIOR APPLICATION NUMBER: 60/164,973
PRIOR FILING DATE: 1999-11-11
NUMBER OF SEQ. ID NOS: 140981
SEQ. ID NOS: 440981
SEQ. ID NO 94301
LENGTH: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Brown, Martin, Haller & McClain
STREET: 1660 Union Street
CITY: San Diego
STATE: California
COUNTRY: USA
ZIP: 91201-2926
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                            Ouery Match 68.0%; Score 13.5; DB 26; Best Local Similarity 80.0%; Pred. No. 1.9e+04; Matches 16; Conservative 0; Mismatches 4;
                                                                                                                                                                                                                                                                                         ; PUBLICATION INFORMATION:
; DATABASE ACCESSION NUMBER: GenBank U51003
US-09-660-220-94301
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CLASSIFCATION NUMBER: USCASS,771A
FILING DATE: September 13, 1994
CLASSIFCATION 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/297,961
FILING DATE: 29-40G-1994
CLASSIFTCATION: 435
PRIOR APPLICATION BATA:
APPLICATION NUMBER: 08/213,446
FILING DATE: 15-MAR-1994
CLASSIFTCATION: 435
PRIOR APPLICATION DATA:
FILING DATE: 15-MAR-1994
CLASSIFTCATION NUMBER: 08/213,447
FILING DATE: 15-MAR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Seidman, Stephanie L.
REGISTRATION NUMBER: 33,779
REFERENCE/DOCKET NUMBER: 519522
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619)238-0999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 53, Application US/08305771A GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (619)238-0062
INFORMATION FOR SEQ ID NO: 53:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 ctaggggaggctcagtgtgg 20
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NAME: Seidman, Stephanie
                                                                                                                                                                                                                                                                   ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 15
US-08-305-771A-53
                                                                                                                                                                                                                                               TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          qq
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Query Match 66.0%;
Best Local Similarity 83.3%;
Matches 15; Conservative
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 JS-09-724-671-20856/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Bovine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TYPE: DNA
; ORGANISM: Mouse
US-09-724-671-20856
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US-60-252-833-23017
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Sequence 24602, A
Sequence 10394, A
Sequence 41965, A
Sequence 2476, Ap
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20856, A Sequence 5380, Ap Sequence 5580, Ap Sequence 4, Appli Sequence 29, Appli Sequence 1694, Appli Sequence 1694, Ap Sequence 10923, A Sequence 1754, Ap Sequence 174, Appli Sequence 14, Appli Sequence 14, Appli Sequence 18, Appli Sequence 18, Appli Sequence 18, Appli Sequence 18, Appli
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19.290 Million cell updates/sec
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                                                                                                                Search time 2173.58 Seconds
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Sequence 5
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1: /cgn2_6/ptodata/2/pna/PCT_NEW_COMB.seq:*

2: /cgn2_6/ptodata/2/pna/USO6_NEW_COMB.seq:*

3: /cgn2_6/ptodata/2/pna/USO8_NEW_COMB.seq:*

4: /cgn2_6/ptodata/2/pna/USO8_NEW_COMB.seq:*

5: /cgn2_6/ptodata/2/pna/USO8_NEW_COMB.seq:*

7: /cgn2_6/ptodata/2/pna/USO8_NEW_COMB.seq:*

8: /cgn2_6/ptodata/2/pna/USO8_NEW_COMB.seq2:*

8: /cgn2_6/ptodata/2/pna/USO8_NEW_COMB.seq2:*
               GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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US-09-198-452A-5586

US-09-653-971A-4

US-09-653-971A-4

US-09-623-972A-1694

US-09-198-452A-1694

US-09-198-452A-1757

US-09-198-452A-1757

US-09-198-452A-1757

US-09-198-452A-1757

US-00-253-378-10923

US-00-253-378-10923

US-00-253-378-10923

US-00-253-378-10923

US-00-253-378-10923

US-00-253-457-11420

US-00-253-457-11420

US-00-253-378-10394

US-00-253-378-10394

US-00-253-378-10394
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US-09-882-246-27
US-60-253-654-8477
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                                                                                                                                                                                                                                                                                                    1895189 seqs, 1048201267 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SUMMARIES
                                                                                                                2001, 16:55:04
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Maximum Match 100%
Listing first 45 summaries
                                                                                 OM nucleic - nucleic search, using sw model
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Minimum DB seq length: 0 Maximum DB seq length: 50
                                                                                                                October
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Match 1
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ALIGNMENTS

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NUMBER OF SEQUENCES: 64
CORRESPONDENCE ADDRESS:
ADDRESSEE: Swanson & Bratschun, L.L.C.
STREET: 84000 E. Prentice Avenue, Suite 200
CITY: Englewood
STATE: Colorado
                 APPLICANT: TESTA, TANIA TAMSON
TESTA TANIA TAMSON
TITLE OF INVENTION: NOVEL COMPOUNDS
FILE REFERENCE: GP-3021
CURRENT APPLICATION NUMBER: US/09/653,971A
CURRENT APPLICATION NUMBER: US/09/653,971A
CURRENT FILING DATE: 2001-08-08
PRIOR FILING DATE: 1999-09-02
PRIOR FILING DATE: 1999-09-02
PRIOR FILING DATE: 2000-03-17
NUMBER OF SEQ ID NOS: 4
SOFTWARE: PASLESQ for Windows Version 3.0
SEQ ID NO + 22
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.44 MG
                                                                                                                                                                                                                                                                                                                                                                                                                Score 11.8; DB 7;
Pred. No. 5.9e+03;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         KOCH, TAD
RINGQUIST, STEVEN
JENSEN, KIRK
ATKINSON, BRENT
ATKINSON, SYSTEMATIC EVOLUTION
OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM
OPERATING SYSTEM: MS-DOS
SOFTWARE: WORDPERFECT 8.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/882,246
FILLING DATE: 14-Jun-2001
CLASSIFICATION: CURROWN>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: CUNKNOWN>
APPLICATION NUMBER: PCT/US94/10542
FILING DATE: 18 September 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 07/714,131
FILING DATE: 10 JUNE 1991
APPLICATION NUMBER: 07/536,428
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 09/093,293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 17 September 1993
APPLICATION NUMBER: 08/143,564
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 08/123,935
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25 October 1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 6
US-09-882-246-29
Sequence 29, Application US/09882246
GENERAL INFORMATION:
    MURDOCK, PAUL RUSSELL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: GOLD, LARRY WILLIS, MICHAEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      storage
                                                                                                                                                                                                                                                                                                                                                                                                                  59.08;
86.78;
                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 59.0
Best Local Similarity 86.7
Matches 13; Conservative
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21 TGTTGGATCCCAATC 7
                                                                                                                                                                                                                                                                                                                TYPE: DNA ORGANISM: HOMO SAPIENS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                        US-09-653-971A-4
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TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragments
TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, preve
TITLE OF INVENTION: and treatment of infection
TITLE OF INVENTION: and treatment of infection
FILE REPERENCE: 9710-003-999
CURRENT APPLICATION NUMBER: US/09/198,452A
CURRENT FILING DATE: 1998-11-24
NUMBER OF SEQ ID NOS: 6849
LENGTH: 20
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                                                              Length 33;
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TITLE OF INVENTION: Compositions isolated from ovine tissues
TITLE OF INVENTION: and methods for their use.
TITLE REFERENCE: 1057P
CURRENT APPLICATION NUMBER: US/60/253,653
CURRENT FILIG DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 30124
SCPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 5980
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 6; Length 20;
                                                                                                    3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                           Score 12.2; DB 8;
Pred. No. 3.8e+03;
0; / Mismatches 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           60.0%; Score 12; DB 100.0%; Pred. No. 4.5
                                                                                                                                                                                                                                                                                          Sequence 5586, Application US/09198452A GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 5980, Application US/60253653 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; TYPE: DNA
; ORGANISM: Chlamydia pneumoniae
US-09-198-452A-5586
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 tgttggtttccaatcggacc 20
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82.4%;
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                                                           Query Match 61.0
Best Local Similarity 82.4
Matches 14; Conservative
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Best Local Similarity 75.0
Matches 15; Conservative
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Matches 12; Conservative
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US-09-198-452A-5586/C
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; ORGANISM: Ovine
US-60-253-653-5980
US-60-252-833-23017
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US-60-253-653-5980
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US-09-653-971A-4/C
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Gaps

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Length 22, Indels us-09-757-100b-15.szlim50.rnpn

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; ORGANISM: Pinus radiata
US-60-253-456-27270
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Best Local Similarity
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US-60-253-378-10923
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: University of Nevada-Reno, Richard Bjur, PhD, JD
TITLE OF INVENTION: Identification of Oxidant Isoform of Human MnSOD
FILE REFERENCE: unevadareno5013
CURRENT APPLICATION NUMBER: US/09/623,025A
CURRENT FILING DATE: 2001-01-16
PRIOR APPLICATION NUMBER: PCT/US99/04129
PRIOR FILING DATE: 1998-02-25
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin Ver. 2.0
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Sequence 27270, Application US/60253456

GENERAL INFORMATION:

APPLICANT: Havukkala, Ilkka J

TITLE OF INVENTION: Polynucleotides, isolated from

CURRENT APPLICATION: DATE: 3060/253,456

CURRENT FILING DATE: 2000-11-27

NUMBER OF SEX ID NOS: 37096

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 27270
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Pred. No. 6.4e+03;
7; Mismatches 2;
                                                                          NAME: Barry J. Swanson
REGISTRATION NUMBER: 33,215
REFERENCE/DOCKET NUMBER: NEX10/US-CON
TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 793-3333
TELEPHONE: (303) 793-3433
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
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0; Mismatches
                                                                                                                                                                                                                                                                                  ; TOPOLOGY: linear ; SEQUENCE DESCRIPTION: SEQ ID NO: 29: US-09-882-246-29
FILING DATE: 11 JUNE 1990
APPLICATION NUMBER: 07/931,
FILING DATE: 17 AUGUST 1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                  LENGTH: 35 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                59.0%;
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Best Local Similarity 92.3
Matches 12; Conservative
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Best Local Similarity
Matches 6; Conserv
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Sequence 1694, Application US/09198452A
GENERAL INFORMATION:
APPLICANT: Griffais, R.
TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, pr
TITLE OF INVENTION: and treatment of infection
FILE REFERENCE: 9710-03-999
CURRENT APPLICATION NUMBER: US/09/198,452A
CURRENT FILING DATE: 1998-11-24
NUMBER OF SEQ ID NOS: 6849
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 4752, Application US/09198452A
GENERAL INFORMATION:
APPLICANT: GIIffals, R.
TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragme;
TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, pr
TITLE OF INVENTION: and treatment of infection
FILE REFERENCE: 9710-003-999/
CURRENT APPLICATION NUMBER: US/09/198,452A
CURRENT FILING DATE: 1998-11-24
NUMBER OF SEQ ID NOS: 6849
SEQ ID NO 4752
LENGTH: 20
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Pred. No. 1.3e+04;
0; Mismatches 3;
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Pred. No. 1.3e+04;
0; Mismatches 3;
        Score 11.4; DB 8;
Pred. No. 1.1e+04;
0; Mismatches 1;
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; ORGANISM: Chlamydia pneumoniae
US-09-198-452A-1694
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; TYPE: DNA
; ORGANISM: Chlamydia pneumoniae
US-09-198-452A-4752
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81.2%;
        57.0%;
92.3%;
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81.2%;
Ouery Match
Best Local Similarity 92.3
Matches 12; Conservative
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Matches 13; Conservative
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Gaps

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Indels

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OTHER INFORMATION: Description of Artificial Sequence: Synthetic DNA
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GENERAL INFORMATION:
APPLICANT: Millenium pharmaceuticals, Inc.
APPLICANT: Regents of the University of California
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING
TITLE OF INVENTION: AND TREATING CHROMOSOME-18p RELATED DISORDERS
FILE REPERENCE: 7853-138-238
CURRENT APPLICATION NUMBER: PCT/US99/05606
CURRENT FILING DATE: 1999-03-16
PRIOR FILING DATE: 1999-01-22
PRIOR FILING DATE: 1999-01-22
NUMBER OF SEQ ID NOS: 84
SOFTWARE: FastSEQ for Windows Version 3.0
; OTHER INFORMATION: of PH1 Fab antibody with additional linker and ; OTHER INFORMATION: restriction sites US-09-822-698A-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
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TITLE OF INVENTION: 19A NEPHROPATHY-ASSOCIATED GENE
FILE REFERENCE: 766.21 CTP
CURRENT APPLICATION NUMBER: US/09/730,559B
CURRENT FILING DATE: 2000-12-07
NUMBER OF SEQ ID NOS: 121
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 55
LENGTH: 24
                                                                                                                                                         Score 11.2; DB 7;
Pred. No. 1.5e+04;
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Pred. No. 1.6e+04;
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NAKAGAWA, SATOSHI
NISHI, TATSUNARI
KUGA, TETSURO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                         56.0%;
81.2%;
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Best Local Similarity 100.
Matches 11; Conservative
                                                                                                                                                                                                                        13; Conservative
                                                                                                                                                                                                                                                                                     2 gttggtttccaatcgg 17
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Best Local Similarity
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US-09-730-559B-55
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PCT-US99-05606-8
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APPLICANT:
APPLICANT:
APPLICANT:
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US-09-822-698A-14/C
Sequence 14, Application US/09822698A
Sequence 14, Application US/09822698A
SEQUENCE 14, Application US/09822698A
SEQUENCE 11 FORMATION:
APPLICANT: Hongenboom, Hendricus R.J.M.
APPLICANT: Hongenboom, Hendricus R.J.M.
TILLE OF INVENTION: MUCIN-1 Specific Binding Members and Methods of Use Thereof
CURRENT APPLICATION NUMBER: US/09/822,698A
CURRENT APPLICATION NUMBER: US 09/538,913
FRIOR APPLICATION NUMBER: US 09/538,913
FRIOR APPLICATION NUMBER: US 09/538,913
SEQUENCE: MICROSOft Word
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GENERAL INFORMATION:
APPLICANT: Glenn, Mattew
APPLICANT: Glenn, Mattew
TITLE OF INVERTION: Compositions isolated from bovine tissue
TITLE OF INVERTION: Compositions of their use.
FILE REFERENCE: 1055P3
CURRENT APPLICATION NUMBER: US/60/253,651
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 27858
SOFTWARE: FastSEQ for Windows Version 4.0
LENGTH: 40
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Pred. No. 1.4e+04;
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1,4e+04;
... 3; Indels
                      TITLE OF INVENTION: Polynucleotides isolated from plants TITLE OF INVENTION: and methods for their use.

TITLE OF INVENTION: and methods for their use.

TITLE OF INVENTION: and methods for their use.

CURRENT APPLICATION NUMBER: US/60/253,378

CURRENT FILING DATE: 2000-11-27

NUMBER OF SEQ ID NOS: 40367

SOFTWARE PASTSEQ for Windows Version 4.0

SEQ ID NO 10923

LENGTH: 38
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Pred. No. 1.4e+04;
); Mismatches 3;
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81.2%;
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81.2%;
   Havukkala, Ilkka J
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 56.0 Best Local Similarity 81.2 Matches 13; Conservative
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Best Local Similarity 81.2
Matches 13; Conservative
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; ORGANISM: Pinus radiata
US-60-253-378-10923
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   APPLICANT:
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Gaps

0; Gaps Query Match 55.0%; Score 11; DB 7; Length 24; Best Local Similarity 73.7%; Pred. No. 1.7e+04; Matches 14; Conservative 0; Mismatches 5; Indels

; 0

1 tgttggtttccaatcggac 19 || || || || || || || || 3 tgatgctttctatctgac 21

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Search completed: October 2, 2001, 16:55:04 Job time: 17667 sec

Run on:

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12.8
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16.701 Million cell updates/sec
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cgn2_6/ptodata/1/pna/US087_COMB.seq:*

cgn2_6/ptodata/1/pna/US088_COMB.seq:*

cgn2_6/ptodata/1/pna/US089_COMB.seq:*

cgn2_6/ptodata/1/pna/US091_COMB.seq:*

cgn2_6/ptodata/1/pna/US092_COMB.seq:*

cgn2_6/ptodata/1/pna/US093_COMB.seq:*

cgn2_6/ptodata/1/pna/US095_COMB.seq:*

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cgn2_6/ptodata/1/pna/US096_COMB.seq:*

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cgn2_6/ptodata/1/pna/US096_COMB.seq:*

cgn2_6/ptodata/1/pna/US098_COMB.seq:*

cgn2_6/ptodata/1/pna/US098_COMB.seq:*
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1: /cgn2_6/ptodata/1/pna/VGO6_COMB.seq:*
2: /cgn2_6/ptodata/1/pna/VGO6_COMB.seq:*
3: /cgn2_6/ptodata/1/pna/VGO0_COMB.seq:*
4: /cgn2_6/ptodata/1/pna/VGO8_COMB.seq:*
5: /cgn2_6/ptodata/1/pna/VGO8_COMB.seq:*
6: /cgn2_6/ptodata/1/pna/VGO8_COMB.seq:*
7: /cgn2_6/ptodata/1/pna/VGO8_COMB.seq:*
8: /cgn2_6/ptodata/1/pna/VGO8_COMB.seq:*
9: /cgn2_6/ptodata/1/pna/VGO8_COMB.seq:*
10: /cgn2_6/ptodata/1/pna/VGO8_COMB.seq:*
11: /cgn2_6/ptodata/1/pna/VGO8_COMB.seq:*
11: /cgn2_6/ptodata/1/pna/VGO8_COMB.seq:*
11: /cgn2_6/ptodata/1/pna/VGO8_COMB.seq:*
11: /cgn2_6/ptodata/1/pna/VGO8_COMB.seq:*
11: /cgn2_6/ptodata/1/pna/VGO8_COMB.seq:*
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6/ptodata/1/pna/US6004_COMB.seq:*
6/ptodata/1/pna/US6005_COMB.seq:*
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/cgn2_6/ptodata/1/pna/US6008_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6009_COMB.seq:*
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/cgn2_6/ptodata/1/pna/US6011_COMB.seq:
GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14155048 seqs, 7388405095 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hits satisfying chosen parameters:
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Maximum Match 100%
Listing first 45 summaries
                                                                                                          OM nucleic - nucleic search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                            IDENTITY_NUC Gaport 1.0
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20
1 tgttggtttccaatcggacc
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Database

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/ Cgn2_6/ptodata/1/pna/US6012_COMB.seq:
/ Cgn2_6/ptodata/1/pna/US6013_COMB.seq:
/ Cgn2_6/ptodata/1/pna/US6014_COMB.seq:
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/ Cgn2_6/ptodata/1/pna/US6019_COMB.seq:
/ Cgn2_6/ptodata/1/pna/US6019_COMB.seq:
/ Cgn2_6/ptodata/1/pna/US6021_COMB.seq:
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/ Cgn2_6/ptodata/1/pna/US6022_COMB.seq:
/ Cgn2_6/ptodata/1/pna/US6028_COMB.seq:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

		Description	Sequence 15	Sequence 15,	-15 Sequence 15,	54082 Sequence 1540	572 Sequence 7572	201094 Sequence	35 Sequence 35, A	35 Sequence 35,	-35 Sequence 35, Ap	1645 Sequence 1645	12731 Sequence 42731,	735 Sednence	Sequence	5701 Sequence	3395/ Sequence	79738 Sedneuce		13724	184627 Sequence	Sequence 6, App	-6 Sequence 6,	261020 Sequence 26	Sednence 69	Sequence 13164	Sequence 13165	Sequence	Sequence 40842	67542 Sequence 16754	09729 Sequence 2	75344 Sequence 27534	76537 Sequence	nonn Soundan	652	122282 Sequence 122282	218413 Sequence	218414 Sequence	5568 Sequenc	17795 Sequence	17795 Sequence	7795 Sequence 17795,	4480 Sequence 3448	84625 Sequence 846
SUMMARIES		B ID	PCT-US00-18999	7 US-09-377-310	001-121-100 001-121-100	5 US-60-234-017	5 US-60-233-620	5 US-60-234-017	PCT-US00-18999	7 US-09-377-3	9 US-09-757-100E	5 US-60-233-620-	5 US-60-234-017-	5 US-60-234-017-	5 US-60-234-017-	5 US-60-234-017-	5 05-60-232-638-	0 US-60-733-T66-	-02C-039-60-310-9	-077-000-60-60 -077-000-63-00 -071-60-530-638-	5 US-60-233-166	3 US-08-983-474-	3 US-08-983	5 US-60-234-017	5 US-60-232-638	5 US-60-232-638	5 US-60-232-638	5 US-60-233-166	5 US-60-233-166	5 US-60-234-017	5 US-60-234-017	S US-60-234-01	55 US-60-234-017-4	0.05-60-634-04	0 03-06-024-033	5 US-60-233-620	5 US-60-234	5 US-60-234-017	8 US-09-422-97	8 US-09-703-70	8 US-60-164-320	0 US-60-183-791	6 US-09-660-220	
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                                                                                                                                                  APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
APPLICANT: Moro, Pamela S.
TITLE OF INVENTION: Expression
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0476
CURRENT APPLICATION NUMBER: PCT/US00/18999
CURRENT FILING DATE: 1999-08-19
FRIOR FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
ATILE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310A
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SEQ ID NO 15
LENCTH: 20
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ilarity 100.0%; Pred. No. 2.5;
Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 20;
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ALIGNMENTS
                                                                                                                                          Isis Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: antisense sequence PCT-US00-18999-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: antisense sequence US-09-377-310-15
                                                                                          Sequence 15, Application PC/TUS0018999 GENERAL INFORMATION:
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Best Local Similarity 100.0
Matches 20; Conservative
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Matches 20; Conserv
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US-09-377-310-15
                                                                                                                                          APPLICANT:
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RESULT 3 US-09-757-100B-15 ; Sequence 15, Application US/09757100B

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APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase;
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0533
CURRENT APPLICATION NUMBER: US/09/757,100B
PRIOR APPLICATION NUMBER: 09/377,310
PRIOR FILING DATE: 1999-08-19
PRIOR FILING DATE: 1999-08-19
PRIOR FILING DATE: 2000-07-13
NUMBER OF SEQ ID NOS: 44
SOFTWARE: PATCHIN VET. 2.0
LENGTH: 20
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APPLICANT: Affymetrix, Inc.
ATTLE OF INVENTION: Methods of Genetic Analysis of Mus
TITLE OF INVENTION: Methods of Genetic Analysis of Mus
FILE REFERENCE: 3115
CURRENT APPLICATION NUMBER: US/60/234,017
CURRENT FILING DATE: 2000-09-20
NUMBER OF SEQ ID NOS: 605887
SOFTWARE: FastSEQ for Windows Version 4.0
ERG ID NO 154082
LENGTH: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 7572, Application US/60233620
GENERAL INFORMATION:
APPLICANT: Mithmann
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of
TITLE OF INVENTION: Arabidopsis thaliana
FILE REFERENCE: 3116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;
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89.5%; Pred. No. 4.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 20; DB; 100.0%; Pred. No. 2.5;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; PUBLICATION INFORMATION:
; DATABASE ACCESSION NUMBER: GenBank A1854607
US-60-234-017-154082
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: antisense sequence US-09-757-100B-15
                                                                                                                                                                                                                                                                                                                                       TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 tgttggtttccaatcggacc 20
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Best Local Similarity 100.
Matches 20; Conservative
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ORGANISM: Mus musculus
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US-60-233-620-7572/c
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Gaps

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Sequence 35, Application US/09377310A

Sequence 35, Application US/09377310A

GENERAL INFORMATION:
APPLICANT: General William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH -0389
CURRENT APPLICATION NUMBER: US/09/377,310A
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 35
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
APPLICANT: Garde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0533
CURRENT APPLICATION NUMBER: US/09/757,100B
CURRENT FILING DATE: 2001-03-15
PRIOR APPLICATION NUMBER: 09/377,310
PRIOR FILING DATE: 1999-08-19
PRIOR APPLICATION NUMBER: PCT/US00/18999
PRIOR APPLICATION NUMBER: PCT/US00/18999
PRIOR APPLICATION NUMBER: PCT/US00/18999
SPRIOR FILING DATE: 2000-07-13
NUMBER OF SEO ID NOS: 44
SOFTWARE: Patentin Ver. 2.0
SEO ID NO 35
LENGTH: 15
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                                                                                                                                       Length 15;
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                                                                                                                                     75.0%; Score 15; DB 1; L 100.0%; Pred. No. 1.2e+03; iive 0; Mismatches 0;
                                               ; OTHER INFORMATION: antisense sequence PCT-US00-18999-35
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; GENERAL INFORMATION:
; APPLICANT: Monia, Brett P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA ORGANISM: Artificial Sequence
    ORGANISM: Artificial Sequence
                                                                                                                                       Query Match 75.0
Best Local Similarity 100.
Matches 15; Conservative
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Best Local Similarity 100.
Matches 15; Conservative
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US-09-377-310-35
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APPLICANT: State P. APPLICANT: Monia, Brett P. APPLICANT: Monia, Brett P. APPLICANT: Gaarde, William A. APPLICANT: Or INVENTION: Antisense Modulation of Focal Adhesion Kinase TITLE OF INVENTION: Expression
FILE REPERENCE: ISPH-0476
CURRENT APPLICATION NUMBER: PCT/USO0/18999
CURRENT FILING DATE: 2000-07-13
PRIOR PRILING DATE: 1999-08-19
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APPLICANT: Mittmann, M.
APPLICANT: Affymetrix, Inc.
TILE OF INVENTION: Methods of Genetic Analysis of Mus; TILE OF INVENTION: Methods of Genetic Analysis of Mus; TILE REPRENCE: 3115
CURRENT APPLICATION NUMBER: US/60/234,017
CURRENT FILING DATE: 2000-09-20
NUMBER OF SEQ ID NOS: 605887
SEQ ID NO 201094
LENGTH: 25
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                                                                                                                                                                                                                                                                                             76.0%; Score 15.2; DB 55;
85.0%; Pred. No. 9.8e+02;
tive 0; Mismatches 3;
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                                                                                                                                                                             ; PUBLICATION INFORMATION:
; DATABASE ACCESSION NUMBER: GenBank AC003028
US-60-233-620-7572
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DATABASE ACCESSION NUMBER: Genbank AA792731
US-60-234-017-201094
CURRENT APPLICATION NUMBER: US/60/233,620
CURRENT FILING DATE: 2000-10-24
NUMBER OF SEQ ID NOS: 131820
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7572
LENGTH: 25
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                                                                                                                                       TYPE: DNA ORGANISM: Arabidopsis thaliana
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SEQ ID NO 35
LENGTH: 15
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Sequence 42735, Application US/60234017 GENERAL INFORMATION:
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   Length 15;
                                    Indels
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GENERAL INFORMATION:
APPLICANT: Mittmann, M
APPLICANT: Affmetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of Mus
TITLE OF INVENTION: Musculus
FITLE REFERENCE: 3115
CURRENT APPLICATION UNBER: US/60/234,017
CURRENT FILING DATE: 2000-09-20
NUMBER OF SEQ ID NOS: 605887
SEQ ID NO 42731
LENGTH: 25
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Sequence 1645, Application US/60233620
Sequence 1645, Application US/60233620
GENERAL INFORMATION:
APPLICANT: Mittmann
APPLICANT: Mittmann
TITLE OF INVENTION: Methods of Genetic Analysis of
TITLE OF INVENTION: Methods of Genetic Analysis of
TITLE OF INVENTION: Arabidopsis thaliana
FILE REFERENCE: 3116
CURRENT APPLICATION NUMBER: US/60/233,620
CURRENT APPLICATION NUMBER: US/60/233,620
CURRENT FILING DATE: 2000-10-24
NUMBER OF SEQ ID NOS: 131820
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1645
LENTH: 25
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Pred. No. 3.4e+03;
0; Mismatches 3;
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84.2%; Pred. No. 3.4e+03;
Live 0; Mismatches 3;
DB 29; L
1.2e+03;
                                      Mismatches
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; DATABASE ACCESSION NUMBER: Genbank AC000132
US-60-233-620-1645
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PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: GenBank L03215
75.0%; Score 15;
100.0%; Pred. No.
:ive 0; Mismatch
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84.28;
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Best Local Similarity 84.2°
Matches 16; Conservative
Query Match 75.(
Best Local Similarity 100.
Matches 15; Conservative
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Best Local Similarity
Matches 16; Conserv
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US-60-234-017-42735/c
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GENERAL INFORMATION:
APPLICANT: Mittmann, M
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of Mus
FILE REFERENCE: 3115
CURRENT APPLICANT: 2000-09-20
NUMBER OF SEQ ID NOS: 605887
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 196005
LENGTH: 25
APPLICANT: Mittmann, M
APPLICANT: Affymetrix, Inc.
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of Mus
TITLE OF INVENTION: Musculus
FILE REFERENCE: 3115
CURRENT APPLICATION NUMBER: US/60/234,017
CURRENT FILING DATE: 2000-09-20
NUMBER OF SEQ ID NOS: 605887
SOFTWARE: FastSEQ for Windows Version 4.0
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GENERAL INFORMATION:
APPLICANT: Mittmann, M
APPLICANT: Affinetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of Mus;
TITLE OF INVENTION: musculus
FILE REFERENCE: 3115;
CURRENT APPLICATION NUMBER: US/60/234,017
CURRENT FILING DATE: 2000-09-20
NUMBER OF SED ID NOS: 605887
SOFTWARE: FastSEQ for Windows Version 4.0
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Pred. No. 3.4e+03;
0; Mismatches 3;
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84.2%; Pred. No. 3.4e+03;
tive 0; Mismatches 3;
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PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank AA175641
US-60-234-017-196005
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DATABASE ACCESSION NUMBER: Genbank L03215

US-60-234-017-42735
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84.2%;
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Best Local Similarity 84.2*
Matches 16; Conservative
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Matches 16; Conservative
                                                                                                                                                                                                                                                                           ORGANISM: Mus musculus
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US-60-234-017-196005/c
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US-60-234-017-216701
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LENGTH: 25
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                                                                                                                                                                                Length 25;
                                                                                                                                                                                                                                                                                                                                                     RESULT 15
US-60-232-638-33957/c
; Sequence 33957, Application US/60232638
; GENERAL INFORMATION:
; APPLICANT: Mittmann
; APPLICANT: Mittmann
; TILE OF INVENTION: Methods of Genetic Analysis of Yeast
TILE OF INVENTION: Methods of Genetic Analysis of Yeast
FILE REPERENCE: 3110
CURRENT PILIKG DATE: 2000-09-14
; WUMBER OF SEC ID NOS: 138410
; SEQ ID NOS: 138410
; SEQ ID NOS: 138410
; SEQ ID NOS: 138410
; SEPE: DNA
; ORGANISM: SACCATOMYCES CETEVISIAE
; PUBLICATION INFORMATION:
; DATABASE ACCESSION NUMBER: SGD YBROOIC
US-60-232-638-33957
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                                     TYPE: DNA
ORGANISM: MUS musculus
ORGANISM: MUSCULUS
DATABASE ACCESSION NUMBER: Genbank AI789319
US-60-234-017-216701
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1 gttggtttacgatcggtcc 19
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Best Local Similarity 88.2*
Matches 15; Conservative
SEQ ID NO 216701
LENGTH: 25
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Search completed: October 2, 2001, 21:50:16 Job time: 24529 sec

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Sequence Sequence Sequence Sequence

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Sequence 151, Application US/09672217
GENERAL INFORMATION:
APPLICANT: FELL, JACK
APPLICANT: DIAZ, MARA
APPLICANT: MCCABE, MERDE
TITLE OF INVENTION: METHOD OF IDENTIFYING PATHOGENIC CRYPTOCOCCI
FILE REFERENCE: 086222/0143
                US-09-371-772B-8141
US-09-371-772B-9045
US-09-371-772B-10818
US-09-371-772B-10817
US-09-371-772B-11311
US-09-371-772B-11314
US-09-371-772B-11354
US-09-371-772B-12297
US-09-371-772B-12297
US-09-371-772B-1335
US-09-708-690-1018
US-09-708-690-11922
US-09-708-690-11922
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US-09-708-690-11922
US-09-708-690-11305
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US-09-708-690-13154
US-09-708-690-1358
US-09-708-690-13641
US-09-708-690-14574
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Pred. No. 4.4e+03;
0; Mismatches 5;
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APPLICANT: Hudson, Thomas J.
APPLICANT: Hudson, Thomas J.
APPLICANT: Lander, Elic S.
APPLICANT: Lander, Elic S.
APPLICANT: Rioux, John
APPLICANT: Siminovitch, Kathy
TITLE OF INVENTION: IBD-RELATED POLYMORPHISMS
FILE REFERENCE: 2825.1025-002
CURRENT APPLICATION NUMBER: US/09/735,271
CURRENT FILING DATE: 2000-12-11
PRIOR APPLICATION NUMBER: US 60/170,257
PRIOR FILING DATE: 1999-12-10
PRIOR FILING DATE: 2000-04-10
PRIOR FILING DATE: 2000-04-10
NUMBER OF SEQ ID NOS: 2058
SOFTWARE: FASTSEQ for Windows Version 4.0
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; OTHER INFORMATION: n = A,T,C or G
US-09-735-271-1541
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75.0%;
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Best Local Similarity 75.0
Matches 15; Conservative
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US-09-735-271-1541/c
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LENGTH: 32
TYPE: DNA
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339, App
339, App
504, App
7196, Ap
9473, Ap
12367, A
12367, A
14371, App
61, App
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3, Appli
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                                                                                                                                    Search time 2173.58 Seconds (without alignments) 19.290 Million cell updates/sec
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1: /cgn2_6/ptodata/2/pna/PCT_NEW_COMB.seq:*

2: /cgn2_6/ptodata/2/pna/USO6_NEW_COMB.seq:*

3: /cgn2_6/ptodata/2/pna/USO8_NEW_COMB.seq:*

4: /cgn2_6/ptodata/2/pna/USO8_NEW_COMB.seq:*

5: /cgn2_6/ptodata/2/pna/USO9_NEW_COMB.seq:*

6: /cgn2_6/ptodata/2/pna/USO9_NEW_COMB.seq:*

7: /cgn2_6/ptodata/2/pna/USO9_NEW_COMB.seq:*

8: /cgn2_6/ptodata/2/pna/USO9_NEW_COMB.seq:*
                  GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   S-09-574-376B-1253
S-09-715-849-504
S-09-7172B-10090
S-09-708-690-9473
S-09-708-690-9473
S-09-708-690-12367
S-09-908-744-61
S-06-253-654-14371
S-09-508-891-1
S-09-908-82-14371
S-09-508-891-1
S-09-508-891-1
S-09-508-891-1
S-09-908-82-27
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US-09-672-217-153

US-09-536-784-339

US-09-765-272-339

US-09-765-272-339

US-09-765-272-339

US-09-716-849-504

US-09-371-7728-10090

US-09-371-7728-10090

US-09-371-7728-10090

US-09-371-7728-1096

US-09-371-7728-14371

US-09-908-744-61

US-09-908-744-61

US-09-908-744-61

US-09-908-744-61

US-09-108-71-1558

US-09-108-81-1

US-09-108-701-1
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Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 1.0
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Match Length
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TITLE OF INVENTION: Streptococcus pneumoniae Antigens and Vaccines
NUMBER OF SEQUENCES: 452
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                     ; OTHER INFORMATION: Description of Artificial Sequence: Primer/Probe US-09-672-217-151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: Description of Artificial Sequence: Primer/Probe US-09-672-217-253
                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: FELL, JACK
APPLICANT: DIA2, MARA
APPLICANT: DIA2, MARA
APPLICANT: MCCABE, MEADE
TITLE OF INVENTION: METHOD OF IDENTIFYING PATHOGENIC CRYPTOCOCCI
FILE REPERENCE: 086222/0143
CURRENT APPLICATION NUMBER: US/09/672,217
CURRENT FILING DATE: 2000-09-29
PRIOR PEDICATION NUMBER: 60/156,598
PRIOR PEDICATION NUMBER: 60/156,598
PRIOR FILING DATE: 1999-09-29
NUMBER OF SEQ ID NOS: 301
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                                                                                                                                                                                                                                                                                                                                        Query Match 63.0%; Score 12.6; DB 7; Length 33; Best Local Similarity 78.9%; Pred. No. 4.4e+03; Matches 15; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 63.0%; Score 12.6; DB 7; Length 33. 78.9%; Pred. No. 4.4e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Human Genome Sciences, Inc. STREET: 9410 Key West Avenue CITY: Rockville STATE: Maryland COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
CURRENT APPLICATION NUMBER: US/09/672,217
CURRENT FILING DATE: 2000-09-29
PRIOR APPLICATION NUMBER: 60/156,598
PRIOR FILING DATE: 1999-09-29
NUMBER OF SEQ ID NOS: 301
SEQ TO NO 151
LENGTH: 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 253, Application US/09672217 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 339, Application US/09536784 GENERAL INFORMATION:
                                                                                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                2 gttagggatggtgccgtca 20
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Matches 15, Conservative
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LENGTH: 33
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Sequence 339, Application US/09765272
GENERAL INFORMATION:
APPLICANT: Choi et. al.
TITLE OF INVENTION: Streptococcus pneumoniae Antigens and Vaccines
NUMBER OF SEQUENCES: 452
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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APPLICATION NUMBER: US/09/765,272 FILING DATE: 22-Jan-2001 CLASSIFICATION: <UNknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 12.6; DB 5;
Pred. No. 4.5e+03;
0; Mismatches 4;
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OPERATING SYSTEM: MSDOS version 6.2
                                    CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/536,784
FILING DATE: 30-Oct-1997
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                   NAME: Michelle S. Marks
REGISTRATION NUMBER: 41,971
REFERENCE/DOCKET NUMBER: PB340P3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: BIOOKES, A. Anders
REGISTRATION NUMBER: 36,373
REFERENCE/DOCKET NUMBER: PB340P2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; SEQUENCE DESCRIPTION: SEQ ID NO: 339: US-09-536-784-339
                                                                                                                                                 APPLICATION NUMBER: 08/961,083
FILING DATE: OCT-30-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA: APPLICATION NUMBER: 08/961,083
                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8504
TELEFAX: (301) 309-8512
INFORMATION FOR SEQ ID NO: 339:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8504
TELEFAX: (301) 309-8512
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: <Unknown>
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                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 37 base pairs
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LENGTH: 37 base pairs
TYPE: nucleic acid
                      SOFTWARE: ASCII Text
                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (301) 309-8 INFORMATION FOR SEQ ID NO: 339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      63.0%;
78.9%;
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Best Local Similarity 78.99
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US-09-765-272-339/c
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                                                                                                        Length 37;
                                                                                                                                               4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: CATAILLY, Michele
APPLICANT: Ireland, James S.
APPLICANT: Lander. Eric S.
TITLE OF INVENTION: HUMAN SINGLE NUCLEOTIDE POLYMORPHISMS
TITLE REFERENCE: 2825.2002-001
CURRENT APPLICATION NUMBER: US/09/715,849
CURRENT FILING DATE: 2000-11-17
PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 589
SOFTWARE: FASESO for Windows Version 4.0
                                                                                                                                                                                                                                                                                                            Sequence 1253, Application US/09574376B
GENERAL INFORMATION:
APPLICANT: Warrington, Janet
APPLICANT: Shah, Nila
APPLICANT: Gingeras, Thomas Raymond
APPLICANT: Application, June
APPLICANT: Gingeras, Thomas Raymond
APPLICANT: Gingeras, Thomas Raymond
TTILE REFERENCE: 3229.2
CURRENT APPLICATION NUMBER: US/09/574,376B
CURRENT FILING DATE: 2000-05-19
NUMBER OF SEQ ID NOS: 1330
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
61.0%; Score 12.2; DB 6;
Best Local Similarity 73.7%; Pred. No. 6.8e+03;
Matches 14; Conservative 1; Mismatches 4;
                                                                                                          Score 12.6; DB 6;
Pred. No. 4.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 12.4; DB 5;
Pred. No. 5.6e+03;
1; Mismatches 2;
                                                                                                                                               0; Mismatches
 TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 339: US-09-765-272-339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-715-849-504/c; Sequence 504, Application US/09715849; GENERAL INFORMATION:
STRANDEDNESS: double
                                                                                                          63.0%;
78.9%;
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81.2%;
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                                                                                                          Query Match 63.0%
Best Local Similarity 78.9%
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13; Conservative
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US-09-574-376B-1253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                             RESULT 6
US-09-574-376B-1253/c
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US-09-715-849-504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 1253
LENGTH: 31
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r.engTH: 21
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APPLICANT: GUNCOUND JUM
APPLICANT: StinchComb, Jum
APPLICANT: StinchComb, Dan
APPLICANT: StinchComb, Dan
APPLICANT: Escobedo, Jaime
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00, 876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1995-08-10
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-01-08
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin version 3.0
                                                                                                                    APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Scoredor, Jaime
APPLICANT: Escobedo, Jaime
APPLICANT: Escobedo, Jaime
APPLICANT: Escobedo, Jaime
APPLICANT: Escobedo, Jaime
APPLICANTON: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Method and Reagent for Vascular Endothelial Growth Factor Receptor
CURRENT APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SSOFTARE: PatentIn version 3.0

LENGTH: 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Description of Artificial Sequence: Enzymatic Nucleic Acid
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                                                           APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Pavco, Pam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 10090, Application US/09371772B GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
Sequence 7196, Application US/09371772B GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Enzymatic Nucleic Acid
US-09-708-690-12367
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GENERAL INFORMATION:
APPLICANT: Gearing, David P.
APPLICANT: Gearing, David P.
TITLE OF INVENTION: HUMAN MICROVASCULAR ENDOTHELIAL LIBRARY
FITLE OF INVENTION: HUMAN MICROVASCULAR ENDOTHELIAL LIBRARY
FILE REFERENCE: 1600.1010-00.2
CURRENT APPLICATION NUMBER: US/09/840,424
CURRENT FILING DATE: 2001-04-23
FRIOR APPLICATION NUMBER: US 60/084,098
PRIOR FILING DATE: 1999-05-04
PRIOR PILING DATE: 1999-03-08
PRIOR PAPLICATION NUMBER: US 60/126,974
PRIOR APPLICATION NUMBER: US 60/126,974
PRIOR PILING DATE: 1999-03-03
PRIOR PILING DATE: 1999-03-04
NUMBER OF SEQ ID NOS: 1311
SOFTWARE: FastSEQ for Windows Version 4.0
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Pred. No. 1.2e+04;
0; Mismatches 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; LOCATION: (1)...(31)
; OTHER INFORMATION: n = A,T,C or G
US-09-840-424-947
                          LENGTH: 38
TYPE: RNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                           Query Match 60.0%;
Best Local Similarity 55.0%;
Matches 11; Conservative
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Richard Peter
Ruchonen Laura
Kolvuranta Kari
Londesborough John
Aristidou Aristos
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81.2%;
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Povelainen Mira
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Best Local Similarity 81.2
Matches 13; Conservative
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  SEQ ID NO 12367
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US-09-908-744-61
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APPLICANT: PRAYCO, Pam
APPLICANT: Stinchcomb, Jam
APPLICANT: Stinchcomb, Jan
APPLICANT: Stinchcomb, Jan
APPLICANT: Stinchcomb, Jan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHBO0,876-L (400/002)
CURRENT APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
PRIOR FILING DATE: 1996-01-08
PRIOR FILING DATE: 1996-01-08
PRIOR FILING DATE: 1996-01-08
PRIOR FILING DATE: 1996-01-01
PRIOR FILING DATE: 2000-10-10
PRIOR FILING DATE: 2000-10-10
PRIOR FILING DATE: 1998-08-10
SRIOR FILING DATE: 1998-08-10
PRIOR FILING DATE: 1999-08-10
PRIOR FILING DATE: 1999-08-10
PRIOR FILING DATE: 3000-10-10
PRIOR FILING DATE: 300-10-10
PRIOR PLING DATE: 300-10-10
PRIOR FILING DATE: 300-10-10
PRIOR PLING DATE: 300-10-10
PRIOR PLING DATE: 300-10-10
PRIOR PLING DATE: 300-10-10
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APPLICANT: PACOC, Pam
APPLICANT: PACOC, Pam
APPLICANT: ACSYIGGEN, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHBOO, 876-L (400/002)
CURRENT APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR PALING DATE: 1995-10-26
PRIOR FILING DATE: 1999-08-10
PRIOR FILING DATE: 1999-08-10
PRIOR FILING DATE: 2000-10-10
NUMBER OF SEQ ID NOS: 20828
SOFTWARE: PatentIn version 3.0
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US-09-708-690-9473
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     Indels
  5
  Mismatches
                                                                                                                                                                                                                                                                       APPLICANT: Ribozyme Pharmaceuticals, Inc.
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GENERAL INFORMATION:
APPLICANT: RIDOZYME Pharmaceuticals, Inc.
APPLICANT: PAYCO, Pam
                                                                                                                                                                                      US-09-708-690-9473; Sequence 9473, Application US/09708690; GENERAL INFORMATION:
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                                                                                                4 gguuacugaugaggccguua 23
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  Conservative
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                               TITLE OF INVENTION: Manufacture of Five-Carbon Sugars and Sugar Alcohols
TITLE OF INVENTION: Manufacture of Five-Carbon Sugars and Sugar Alcohols
FILE REPERENCE: 1427.0010005;
CURRENT APPLICATION NUMBER: US/09/908,744
CURRENT FILING DATE: 2001-07-20
PRIOR APPLICATION NUMBER: US 09/488,581
PRIOR FILING DATE: 2000-01-21
PRIOR FILING DATE: 1000-01-21
PRIOR FILING DATE: 1997-01-29
PRIOR FILING DATE: 1997-01-29
PRIOR FILING DATE: 1995-01-29
PRIOR FILING DATE: 1995-01-39
PRIOR FILING DATE: 1995-01-03
PRIOR FILING DATE: 1995-01-03
PRIOR FILING DATE: 1995-01-05
PRIOR FILING DATE: 1992-11-05
NUMBER OF SEQ ID NOS: 70
SOFTWARE: PatentIn Ver. 3.0
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| US-60-253-654-14371/c
| US-60-253-654-14371/c
| US-60-253-654-14371, Application US/60253654
| GENERAL INFORMATION:
| APPLICANT: Glenn, Matthew
| APPLICANT: Glenn, Matthew
| TITLE OF INVENTION: Compositions isolated from forage
| TITLE OF INVENTION: Compositions isolated from forage
| TITLE OF INVENTION: Grasses and methods for their use.
| FILE REPERENCE: 1056P
| CURRENT APPLICATION NUMBER: US/60/253,654
| CURRENT FILING DATE: 2000-11-28
| NUMBER OF SEQ ID NOS: 32581
| SOFTWARE: FastSEQ for Windows Version 4.0
| SEQ ID NO 14371
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Pred. No. 1.2e+04;
0; Mismatches 2;
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Pred. No. 1.
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Penttil, Merja
Plazanet-Menut Claire
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; ORGANISM: Festuca arundinaceae
US-60-253-654-14371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        59.0%;
86.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA ORGANISM: Artificial Sequence
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Best Local Similarity 77.8%;
Matches 14; Conservative
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Best Local Similarity
Matches 13; Conserv
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y APPLICANT: Norriss, Michael G.
TITLE OF INVENTION: Compositions isolated from forage
TITLE OF INVENTION: Compositions isolated from forage
TITLE OF INVENTION: grasses and methods for their use.
TITLE REFERENCE: 1056p2
CURRENT APPLICATION NUMBER: US/60/255,592
CURRENT FILING DATE: 2000-12-14
CURRENT FILING DATE: 2000-12-14
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 14371
TENGTH: 26
TYPE: DNA
CORANISM: Festuca arundinaceae
US-60-255-592-14371
QUETY MATCh
Best Local Similarity 77.8%; Pred. No. 1.4e+04;
Matches 14; Conservative 0; Mismatches 4; Indels 0; Gaps
QY
3 ttagggatggtgcgtca 20
II | | | | | | | | | | |
Db 18 TTTGGATGGTTGGTCA 1
Search completed: October 2, 2001, 16:55:04
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16.701 Million cell updates/sec
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| cgn2_6/ptodata/1/pna/PCTUS_COMB.seq:*
| cgn2_6/ptodata/1/pna/USO6_COMB.seq:*
| cgn2_6/ptodata/1/pna/USO8_COMB.seq:*
| cgn2_6/ptodata/1/pna/USO9_COMB.seq:*
| cgn2_6/ptodata/1/pna/USO9_C
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/cgn2_6/ptodata/1/pna/US6007_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6008_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6009_COMB.seq:*
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/cgn2_6/ptodata/1/pna/US6011_COMB.seq:*
version 4.5
- 2000 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                  - nucleic search, using sw model
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   GenCore
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Perfect score:
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/ Cgn2_6/ptodata/1/pna/US6015_COMB.seq:
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/ Cgn2_6/ptodata/1/pna/US6015_COMB.seq:
/ Cgn2_6/ptodata/1/pna/US6019_COMB.seq:
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/ Cgn2_6/ptodata/1/pna/US6019_COMB.seq:
/ Cgn2_6/ptodata/1/pna/US6020_COMB.seq:
/ Cgn2_6/ptodata/1/pna/US6021_COMB.seq:
/cgn2_6/ptodata/1/pna/Us6012_COMB.seq:*/cgn2_6/ptodata/1/pna/Us6013_COMB.seq:*/cgn2_6/ptodata/1/pna/Us6014_COMB.seq:*/cgn2_6/ptodata/1/pna/Us6014_COMB.seq:*/
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/cgn2_6/ptodata/1/pna/US6028_COMB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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	equence 14, Sequence 14, Sequence 14, Sequence 195	34, Appl 34, App 34, App 13935, 39603,	Sequence 79723, A Sequence 255450, Sequence 58, Appl Sequence 58, Appl Sequence 1, Appl Sequence 18, Appl Sequence 11, Appl Sequence 11, Appl Sequence 118931.		88045 88045 1120, 46 1121, Ap 1120, Ap 1121, 32127 1131, 32127 894150 89155 89155 89155 13840
SUMMAKLES	-18999-1 77-310-1 57-100B- 33-166-1	00-18999-3 -377-310-3 -757-100B- -660-220-1		234-017-472 234-049-141 233-166-628 233-166-285 233-166-285 233-166-285 233-166-285 234-017-336	S. 60 - 234 - 041 - 059
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GENERAL INFORMATION:
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                                                                                                                                                       APPLICANT: Monia, Brett P.

APPLICANT: Gaarde, William A.

APPLICANT: Gaarde, William A.

APPLICANT: Gaarde, William A.

APPLICANT: Nero, Pamela S.

TITLE OF INVENTION: Expression

FILE REFERENCE: ISPH-0476

CURRENT APPLICATION NUMBER: PCT/US00/18999

CURRENT FILING DATE: 2000-07-13

PRIOR FILING DATE: 1999-08-19

NUMBER OF SEQ ID NOS: 43

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 14

LENGTH: 20
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APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
ATILE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310A
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
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Pred. No. 10;
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100.0%; Pred. No. ...
0; Mismatches
               ALIGNMENTS
                                                                                                                                           Isis Pharmaceuticals, Inc.
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                                                                                                  Sequence 14, Application PC/TUS0018999
GENERAL INFORMATION:
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US-09-757-100B-14
; Sequence 14, Application US/09757100B
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Best Local Similarity 100.(
Matches 20; Conservative
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Matches 20; Conservative
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Gaps
APPLICANT: Garade, William A.
APPLICANT: Garade, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0533
CURRENT APPLICATION NUMBER: US/09/775,100B
CURRENT FILING DATE: 2001-03-15
PRIOR APPLICATION NUMBER: PCT/US00/18999
PRIOR PILING DATE: 1999-08-19
PRIOR PILING DATE: 2000-07-13
NUMBER OF SEQ ID NOS: 44
SOFTWARE: PATENTIN Ver. 2.0
LENGTH: 20
LENGTH: 20
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APPLICANT: Gaarde, William A.
APPLICANT: Nero, Pamela S.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
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APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of the Rat
TITLE OF INVENTION: Genome
FILE REFERENCE: 3112
CURRENT APPLICATION NUMBER: US/60/233,166
CURRENT FILING DATE: 2000-10-24
NUMBER OF SEQ ID NOS: 420907
SOFTWARE: FastSEQ for Windows Version 4.0
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85.0%; Pred. No. 2.6e+03;
tive 0; Mismatches 3;
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10;
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; PUBLICATION INFORMATION:
; DATABASE ACCESSION NUMBER: Genbank AA998576
US-60-233-166-198784
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 20; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 198784, Application US/60233166
; GENERAL INFORMATION:
; APPLICANT: Mittmann
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                                                                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: antisense sequence US-09-757-100B-14
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PCT-USO0-18999-34
; Sequence 34, Application PC/TUSO018999
; GRERAL INCRMATION:
; APPLICANT: Isis Pharmaceuticals, Inc
                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
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Matches 17; Conservative
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Best Local Similarity
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LENGTH: 25
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Gaps

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Length 25;
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GENERAL INFORMATION:
APPLICANT: Mittmann et al.
APPLICANT: Affymetrix, Inc.
TITLE OF INVATION: Methods of Genetic Analysis of Human
FILE REFERENCE: 3102.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 39603, Application US/60232638
GENERAL INFORMATION:
APPLICANT: Mittmann
APPLICANT: Affymertix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of Yeast
FILE REFERENCE: 3110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  71.0%; Score 14.2; DB 26;
84.2%; Pred. No. 8.2e+03;
                                                                                                                                                                                                                               Query Match 75.0%; Score 15; DB 29; Best Local Similarity 100.0%; Pred. No. 3.2e+03; Matches 15; Conservative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/660,220 CURRENT FILLING DATE: 2000-09-12 PRIOR APPLICATION NUMBER: 60/164,973 PRIOR FILLING DATE: 1999-11-11 NUMBER OF SEQ ID NOS: 140981 SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 13935
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: 3110
CURRENT APPLICATION NUMBER: US/60/232,638
CURRENT FILING DATE: 2000-09-14
NUMBER OF SEQ ID NOS: 138410
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 39603
LENGTH: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; PUBLICATION INFORMATION:
; DATABASE ACCESSION NUMBER: Genbank D83784
US-09-660-220-13935
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DATABASE ACCESSION NUMBER: SGD YCL029C
                                                                                                                                              ; OTHER INFORMATION: antisense sequence US-09-757-1008-34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA ORGANISM: Saccaromyces cerevisiae
                                                                                                       ORGANISM: Artificial Sequence
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SEQ ID NO 34
LENGTH: 15
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Matches 16; Conservative
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ORGANISM: Homo sapiens
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US-09-660-220-13935
  NUMBER OF SEQ
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                                                                                   TYPE: DNA
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Sequence 34, Application US/09757100B

Sequence 34, Application US/09757100B

GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
APPLICANT: Garde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase;
TITLE OF INVENTION: Expression
FILE OF INVENTION: Expression
FILE OF INVENTION: UNMER: US/09/757,100B

CURRENT FILING DATE: 2001-03-15
PRIOR APPLICATION NUMBER: 09/377,310
PRIOR FILING DATE: 1999-08-19
PRIOR FILING DATE: POT/US00/18999

PRIOR FILING DATE: 2000-07-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310A
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
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             FILE REFERENCE: ISPH-0476
CURRENT APPLICATION NUMBER: PCT/USO0/18999
CURRENT FILING DATE: 2000-07-13
PRIOR APPLICATION NUMBER: 09/377,310
PRIOR FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: PATENTIN VET: 2.0
                                                                                                                                                                                                                                                                     ; OTHER INFORMATION: antisense sequence PCT-US00-18999-34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: antisense sequence US-09-377-310-34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-377-310-34; Sequence 34, Application US/09377310A; GENERAL INFORMATION:
  Expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.
Matches 15; Conservative
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Best Local Similarity 100.
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
                                                                                                                                                              SEQ ID NO 34
LENGTH: 15
                                                                                                                                                                                                            TYPE: DNA
                                                                                                                                                                                                                                                     FEATURE:
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Gaps

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Sequence 58, Application US/09004729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: COUNCIL, GATY J.
REGISTRATION NUMBER: 32,020
REFRENCE/DOCKET NUMBER: 2618
TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 863-9700
INFORMATION FOR SED ID NO: 58:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sheridan Ross P.C.
                                                                                                                                                                                                                                                         ZIP: 80203
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TOPOLOGY: linear
; MOLECULE TYPE: DNA (primer)
US-09-004-729-58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           71.0%;
84.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 34 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Sheridan 1
                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
CLASSIFICATION: 424
                                                                                                                                                                                                                        Colorado
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
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STATE:
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 Length 25;
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Sequence 255450, Application US/60234017

GENERAL INFORMATION:

APPLICANT: Mittmann, M

APPLICANT: Affymetrix, Inc.

TITLE OF INVENTION: Methods of Genetic Analysis of Mus

FILE REFERENCE: 3115

CURRENT APPLICATION NUMBER: US/60/234,017

CURRENT PILING DATE: 2000-09-20

NUMBER OF SEQ ID NOS: 605887

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 255450
                                         Indels
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                                                                                                                                                                                            Sequence 79723. Application US/60233620
GENEBAL INFORMATION:
APPLICANT: Mittmann
TITLE OF INVERTION: Methods of Genetic Analysis of
TITLE OF INVERTION: Methods of Genetic Analysis of
TITLE OF INVERTION: Arabidopsis thaliana
FILLE REFERENCE: 3116
CURRENT APPLICATION NUMBER: US/60/233,620
CURRENT FILING DATE: 2000-10-24
NUMBER OF SEQ ID NOS: 131820
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 79723
71.0%; Score 14.2; DB 55;
11arity 84.2%; Pred. No. 8.2e+03;
Conservative 0; Mismatches 3;
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Pred. No. 8.2e+03;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  71.0%; Score 14.2; DB 55;
84.2%; Pred. No. 8.2e+03;
tive 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; PUBLICATION INFORMATION:
; DATABASE ACCESSION NUMBER: GenBank AC005168
US-60-233-620-79723
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank AW125675
US-60-234-017-255450
                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Arabidopsis thallana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 71.0%;
Best Local Similarity 84.2%;
Matches 16; Conservative
                                                                         1 ggttagggatggtgccgtc 19
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                                                                                            24 GGTCAGCTATGGTGCCGTC 6
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Best Local Similarity 84.2
Matches 16; Conservative
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Query Match
Best Local Similarity
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US-09-004-729-58
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                                   Matches
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APPLICANT: Wu Hunter, Shirley
APPLICANT: Stiegler, Gary
APPLICANT: Stiegler, Gary
APPLICANT: Gaines, Patrick J.
TITLE OF INVENTION: FLEA PROTEASE PROTEINS, NUCLEIC ACID
TITLE OF INVENTION: MOLECULES AND USES THEREOF
NUMBER OF SEQUENCES: 103
CORRESPONDENCE ADDRESS:
ADDRESSE: ADDRESSE: ADDRESSE: Sheridan Ross P.C.
STREET: 1700 Lincoln Street, Suite 3500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 13
US-09-004-755-58
US-09-004-755-58
Sequence SB, Application US/09004755
GENERAL INFORMATION:
APPLICANT: Wu Hunter, Shirley
APPLICANT: Stiegler, Gary
APPLICANT: Gaines, Patrick J.
TITLE OF INVENTION: FLEA PROTEASE PROTEINS, NUCLEIC ACID
TITLE OF INVENTION: MOLECULES AND USES THEREOF
NUMBER OF EXQUENCES: 103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/004,729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 14.2; DB 14;
Pred. No. 8.3e+03;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/749,699
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 80203
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMpatible
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RESULT 15
US-07-896-094-1/c
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APPLICANT: Rushlow, Keith E.
APPLICANT: Rushlow, Keith E.
APPLICANT: Rushlow, Keith E.
APPLICANT: Frank, Glenn R.
APPLICANT: Stiegler, Gary
APPLICANT: Stiegler, Gary
TITLE OF INVENTION: FLEA PROTEASE PROTEINS, NUCLEIC ACID MOLECULES AND USES TITLE OF INVENTION: THEREOF INVENTION: THEREOF INVENTION: 1999-04-23
FILE REFERENCE: 2618-25-C4-PUS
CURRENT APPLICATION NUMBER: 1999-04-23
PRIOR APPLICATION NUMBER: PCT/US97/06121
PRIOR APPLICATION NUMBER: PCT/US97/06121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 14.2; DB 14; Length 34;
Pred. No. 8.3e+03;
0; Mismatches 3; Indels (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: Description of Artificial Sequence:primer US-09-171-741C-58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                  PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 14.2; DB 15;
Pred. No. 8.3e+03;
0; Mismatches 3;
                                                                                        CLASSIFICATION:
PRIOR APPLICATION:
APPLICATION DATA:
APPLICATION NUMBER: US/08/749,699
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Connell, Gary J.
REGISTRATION NUMBER: 32,020
REFERENCE/DOCKET NUMBER: 2618-25-C3
TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 863-9700
TELEFERX: (303) 863-9700
TELEFERX: (303) 863-973
INFORMATION FOR SEQ ID NO: 58:
SEQUENCE CHARACTERISTICS:
LENGTH: 34 base pairs
TYPE: NUCLEIC acid
STRANDENNESS: SINGIE
                                                                                                                                                                                                                                       2618-25-C3
                                 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/004,755
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 58, Application US/09171741C
GENERAL INFORMATION:
APPLICANT: Grieve, Robert B.
APPLICANT: Rushlow, Keith E.
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   71.0%;
84.2%;
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84.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                               ; MOLECULE TYPE: DNA (primer) US-09-004-755-58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 gttagggatggtgccgtca 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13 GTTAGCGATCGTCCCGTCA 31
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Best Local Similarity 84.2
Matches 16; Conservative
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SOFTWARE: PatentIn Ve
SEQ ID NO 58
LENGTH: 34
                                                                                                                                                                                                                                                                                                                                                                                                                linear
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Best Local Similarity
Matches 16; Conserv
                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                ropology:
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                                     APPLICANT: Sheiness, Diana K.
APPLICANT: Adams, Trevor H.
APPLICANT: Stamm, Michael P.
APPLICANT: Cangelosi, Gerard A.
TITLE OF INVENTION: Methods and Pharmaceutical Kits Useful
TITLE OF INVENTION: Infections Microorganisms Associated with Vaginal
TITLE OF INVENTION: Infections
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                   ADDRESSEE: Townsend and Townsend
STREET: One Market Plaza, Steuart Tower, Suite 2000
CITY: San Francisco
STATE: California
ZIP: 94105
                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/896,094
FILING DATE: 19920529
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen L.
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 11652-73-1
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION SEQ 1D NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 13.8; DB 3;
Pred. No. 1.3e+04;
0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
Sequence 1, Application US/07896094 GENERAL INFORMATION:
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88.2%;
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EDNESS: single
                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Search completed: October
Job time: 24528 sec
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Best Local Similarity
Matches 15; Conserv
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                                                                                                    2, 2001, 21:50:14; Search time 17695.9 Seconds (without alignments) 16.701 Million cell updates/sec
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/cgn2_6/ptodata/1/pna/US096C_COMB.seq:*
/cgn2_6/ptodata/1/pna/US096D_COMB.seq:*
/cgn2_6/ptodata/1/pna/US096E_COMB.seq:*
/cgn2_6/ptodata/1/pna/US097A_COMB.seq:*
/cgn2_6/ptodata/1/pna/US097A_COMB.seq:*
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/cgn2_6/ptodata/1/pna/US097C_COMB.seq:*
/cgn2_6/ptodata/1/pna/US098_COMB.seq:*
/cgn2_6/ptodata/1/pna/US098_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6000_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6000_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6001_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6002_COMB.seq:*
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6/ptodata/1/pna/US6005_COMB.seq:*
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/ptodata/1/pna/US6008_COMB.seq:*
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/cgn2_6/ptodata/1/pna/US6011_COMB.seq:*
GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                            14155048 seqs, 7388405095 residues
                                                                                                                                                                                                                                                                                                                                            Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Maximum Match 100%
Listing first 45 summaries
                                                                      OM nucleic - nucleic search, using sw model
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/cgn2_6/ptodate/1/pna/US6012_COMB.seq:*
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/cgn2_6/ptodate/1/pna/US6020_COMB.seq:*
/cgn2_6/ptodate/1/pna/US6020_COMB.seq:*
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/cgn2_6/ptodate/1/pna/US6022_COMB.seq:*
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/cgn2_6/ptodate/1/pna/US6022_COMB.seq:*
/cgn2_6/ptodate/1/pna/US6022_COMB.seq:*
/cgn2_6/ptodate/1/pna/US6025_COMB.seq:*
/cgn2_6/ptodate/1/pna/US6026_COMB.seq:*
/cgn2_6/ptodate/1/pna/US6026_COMB.seq:*
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/cgn2_6/ptodate/1/pna/US6026_COMB.seq:*
/cgn2_6/ptodate/1/pna/US6026_COMB.seq:*
  /cgn2_6/ptodata/1/pna/US6012_COMB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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GENERAL INFORMATION:
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                                                                                                                                                                           APPLICANT: ASIS FINIMACCULICALS, INC.
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
APPLICANT: Garde, William A.
APPLICANT: Moro, Pamela S.
TITLE OF INVENTION: Expression
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0476
CURRENT APPLICATION NUMBER: PCT/USOO/18999
CURRENT APPLICATION NUMBER: 09/377,310
PRIOR APPLICATION NUMBER: 09/377,310
PRIOR FILING DATE: 1999-08-19
WUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Montan, Brett P.
APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310A
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
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                                                                                                                                                              Isis Pharmaceuticals, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: antisense sequence PCT-US00-18999-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CTHER INFORMATION: antisense sequence US-09-377-310-12
                                                                                                                      Sequence 12, Application PC/TUS0018999 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-00-377-310-12; Sequence 12, Application US/09377310A; GENERAL INFORMATION:
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Best Local Similarity 100.0
Matches 20; Conservative
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Best Local Similarity 100.(
Matches 20; Conservative
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-09-757-100B-12
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LENGTH: 20
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LENGTH: 20
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Sequence 16, Application PC/TUS0007678
; Sequence 16, Application PC/TUS0007678
; GENERAL INFORMATION:
    APPLICANT: Word, Alexander K.C.
    APPLICANT: Tavtigian, Sean V.
    APPLICANT: Tavtigian, Sean V.
    APPLICANT: Tavtidian, Sean V.
    APPLICANT: Tavtidian, Sean V.
    APPLICANT: Myriad Genetics, Inc.
    TITLE OF INVENTION: BRG1 IS A TUMOR SUPPRESSOR THAT IS MUTATED IN PROSTATE
    TITLE OF INVENTION: AND OTHER CANCER TYPES
    TITLE OF INVENTION: AND OTHER CANCER TYPES
    TITLE OF INVENTION: AND OTHER CANCER TYPES
    CURRENT APPLICATION NUMBER: PCT/US00/07678
    CURRENT FILING DATE: 1999-03-23
    EARLIER APPLICATION NUMBER: U.S. 60/125,806
    SOFTWARE: PLING DATE: 1999-03-23
    NUMBER OF SEQ ID NOS: 77
    SOFTWARE: PatentIn Ver. 2.0
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APPLICANT: Garde, William A.
APPLICANT: Garde, William A.
APPLICANT: Nero, Pamela S.
TITLE OF INVENTION: Expression
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-053
CURRENT APPLICATION NUMBER: US/09/757,100B
CURRENT APPLICATION NUMBER: US/09/757,100B
PRIOR PILING DATE: 1999-08-19
PRIOR PILING DATE: 1999-08-19
PRIOR FILING DATE: 2000-07-13
NUMBER OF SEQ ID NOS: 44
SOFTWARE: Patentin Ver. 2.0
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Pred. No. 6.1e+02;
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Pred. No. 9.7;
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APPLICANT: Wong, Alexander K.C.
APPLICANT: Tavtiglan, Sean V.
APPLICANT: Teng, David H.-F.
                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
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Best Local Similarity 94.4
Matches 17; Conservative
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PCT-US00-07678-16/c
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LENGTH: 20
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LENGTH: 20
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; PUBLICATION INFORMATION:
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TITLE OF INVENTION: BRG1 IS A TUMOR SUPPRESSOR THAT IS MUTATED IN PROSTATE TITLE OF INVENTION: AND OTHER CANCER TYPES
FILE REPERBORCE: 2318-259
CURRENT APPLICATION NUMBER: US/09/535,008
CURRENT FILING DATE: 2000-03-23
EARLIER APPLICATION NUMBER: U.S. 60/125,806
EARLIER FILING DATE: 1999-03-23
NUMBER OF SEQ ID NOS: 77
SOFTWARE: PATENTIN Ver. 2.0
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; Sequence 196919, Application US/60233166
; GENERAL INFORMATION:
; APPLICANT: Mittmann
; APPLICANT: Mittmann
; TITLE OF INVENTION: Genome
; TITLE OF INVENTION: Genome
; TITLE OF INVENTION: Genome
; CURRENT APPLICATION UNMER: US/60/233,166
; CURRENT FILING DATE: 2000-10-24
; WUMBER OF SEQ ID NOS: 420907
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 196919
                                                                                                                                                                                                                                                                                                                            Length 20;
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GENERAL INFORMATION:
APPLICANT: Mittmann
APPLICANT: Mittmann
APPLICANT: Affirmetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of the Rat
TITLE OF INVENTION: Genome
FILE REFERENCE: 3112
CURRENT APPLICATION NUMBER: US/60/233,166
CURRENT APPLICATION NUMBER: US/60/233,166
SOFTWARE: FastSEQ for Windows Version 4.0
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Pred. No. 6.1e+02;
0; Mismatches 1;
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Pred. No. 1.2e+03;
0; Mismatches 2;
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; DATABASE ACCESSION NUMBER: GenBank AA998576
US-60-233-166-196919
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94.4%;
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Best Local Similarity 89.5%;
Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                20 CTGGCATCAGTAGCATCT 3
                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                             ; ORGANISM: Homo sapiens
US-09-535-008-16
                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 17; Conserv
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LENGTH: 25
                                                                                                                                                                                   SEQ ID NO 16
LENGTH: 20
                                                                                                                                                                                                                            TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Gaarde, William A. APPLICANT: Nero, Pamela S. TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase TITLE OF INVENTION: Expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 32, Application US/09377310A
GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
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                                                                                    Length 25;
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                                                                                                                              Indels
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                                                                                Score 15.8; DB 55;
Pred, No. 1.2e+03;
0; Mismatches 2;
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2.9e+03;
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Live 0; Mismatches
; DATABASE ACCESSION NUMBER: GenBank A1639536
US-60-233-166-296426
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: ISPH-0476
CURRENT APPLICATION NUMBER: PCT/US00/18999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310A
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            UMBER: 09/377,310
1999-08-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: antisense sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: antisense sequence US-09-377-310-32
                                                                                                                                                                                                                                                                                              PCT-US00-18999-32; Sequence 32, Application PC/TUS0018999; GENERAL INFORMATION: APPLICANT: Isis Pharmaceuticals, Inc. APPLICANT: Monia, Brett P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT FILING DATE: 2000-07-13
                                                                                    79.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA ORGANISM: Artificial Sequence
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                                                                                                                                                                                               22 CTGACATAAGTAGCACCTC 4
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SOFTWARE: Patentin Ver. 2.0
                                                                                    Query Match 79.0
Best Local Similarity 89.5
Matches 17; Conservative
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Best Local Similarity 100.
Matches 15; Conservative
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PRIOR FILING DATE: 1999-0
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Matches

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Sequence 198168, Application US/60233166
SENERAL INFORMATION:
APPLICANT: Mittmann
TITLE OF INVENTION: Methods of Genetic Analysis of the Rat
TITLE OF INVENTION: Methods of Genetic Analysis of the Rat
TITLE OF INVENTION: Genome
FILE REFERENCE: 3112
CURRENT APPLICATION NUMBER: US/60/233,166
CURRENT FILIG DATE: 2000-10-24
NUMBER OF SEQ ID NOS: 420907
SEQ ID NO 198168
LENGTH: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 25;
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                                                                                                           Sequence 198167, Application US/60233166
GENERAL INFORMATION:
APPLICANT: Mittmann
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Genome
FILE REFERENCE: 3112
CURRENT APPLICATION UNMBER: US/60/233,166
CURRENT FILING DATE: 2000-10-24
NUMBER OF SEQ ID NOS: 420907
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Mittmann APPLICANT: Affymetrix, Inc. TITLE OF INVENTION: Methods of Genetic Analysis of the Rat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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Pred. No. 7.9e+03;
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84.2%; Pred. No. 7.9e+03;
iive 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                          ; PUBLICATION INFORMATION:
; DATABASE ACCESSION NUMBER: GenBank AA998576
US-60-233-166-198167
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PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank AA998576
                                                                                                                                                                                                                                                                                    SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 198167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Sequence 303481, Application US/60233166; GENERAL INFORMATION:
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84.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 cggacaccagtagcatcac 20
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Best Local Similarity 84.2
Matches 16; Conservative
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                        18 CTGACATCAACAGCATCT
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Best Local Similarity
Matches 16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
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    Gaps
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                                                                                                                                                                                                         APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
APPLICANT: Nero, Pamela S.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REPERRACE: ISPH-053
CURRENT APPLICATION NUMBER: US/09/757,100B
CURRENT FILING DATE: 2001-03-15
PRIOR APPLICATION NUMBER: PCT/US00/18999
PRIOR FILING DATE: 2000-07-13
NUMBER OF SEQ ID NOS: 44
SOFTWARE: PatentIn Ver. 2.0
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    Indels
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Pred. No. 3.8e+03;
0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Brown, Thomas A.
APPLICANT: Do Wet, Jeffrey R.
APPLICANT: Gowen, Lori C.
APPLICANT: Hames, Lynn M.
TITLE OF INVENTION: Mammalian Osteoregulins
FILE REFERENCE: PC10445
CURRENT APPLICATION NUMBER: US/09/794,422
CURRENT FILING DATE: 2001-02-27
PRIOR FILING DATE: 2000-02-29
PRIOR APPLICATION NUMBER: 60/185,617
PRIOR APPLICATION NUMBER: 60/234,500
PRIOR FILING DATE: 2000-09-22
  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: antisense sequence US-09-757-1008-32
                                                                                                                                                  US-09-757-100B-32; Sequence 32, Application US/09757100B; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-794-422-41/c
; Sequence 41, Application US/09794422
; GENERAL INFORMATION:
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88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA ORGANISM: Artificial Sequence
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  Conservative
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Matches 16; Conservative
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Matches 15; Conservative
                                  3 tgacatcagtagcat 17
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; ORGANISM: Homo sapiens
US-09-794-422-41
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SOFTWARE: Patentin Ve
SEQ ID NO 41
LENGTH: 20
15;
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Query Match

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Query Match

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Gaps

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Length 17; Indels

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Score 13.8; DB 8;
Pred. No. 1.2e+04;
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                                                                                                                                                                                                                 69.0%;
88.2%;
  INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 17 base pairs TYPE: nucleic acid STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                           4 gacatcagtagcatctc 20
                                                                                                                                                                                                                                                                Conservative
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Best Local Similarity
Matches 15; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1431, Application US/08435632
GENERAL INFORMATION:
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Draper, Kenneth
APPLICANT: Draper, Kenneth
APPLICANT: McSwiggen, James
APPLICANT: Carver
APPLICANT: Correct Carver
TITLE OF INVENTION: TREATMENT OF RESTENOSIS AND
TITLE OF INVENTION: CANCER USING RIBOZYMES
NUMBER OF SEQUENCES: 2627
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: ...
ZIP: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/435,632
FILLIAG DATE: 05-MAY-1995
FILLIAG DATE: 05-MAY-1995
TITLE OF INVENTION: Genome
FILE REFERENCE: 3112
CURRENT APPLICATION NUMBER: US/60/233,166
CURRENT FILING DATE: 2000-10-24
NUMBER OF SEQ ID NOS: 420907
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 303481
LENGTH: 25
                                                                                                                                                                                           TYPE: DNA
ORGANISM: Rattus norvegicus
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: GenBank D10757
US-60-233-166-303481
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REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 209/035
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFA: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PULLOR DATE:

APPLICATION DATA:

APPLICATION NUMBER: 08/373,124
FILING DATE: January 13, 1995
APPLICATION NUMBER: 08/245,466
FILING DATE: May 18, 1994
APPLICATION NUMBER: 07/192,943
FILING DATE: Pebruary 7, 1994
APPLICATION NUMBER: 07/987,132
FILING DATE: December 7, 1992
APPLICATION NUMBER: 07/96,422
FILING DATE: AUGUST 26,1992
ATOMNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 cctgacatcagtagcatct 19
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STATE: California
COUNTRY: U.S.A.
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Best Local Similarity
Matches 16; Conserv:
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US-08-435-632-1431/c
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Gaps
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Patent No. 6133031
GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase;
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT APPLICATION NUMBER: US/09/377,310B
CURRENT PILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 3
LENGTH: 20
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Sequence
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Sequence
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APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Expression
FILE REPERENCE: ISPH-0389
CURRENT APPLICATION VIMBER: US/09/377,310B
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 23
LENGTH: 15
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        US-08-634-826-8
US-09-115-175-8
US-09-015-175-8
US-08-096-947-7
US-08-096-947-9
US-07-919-140B-9
US-07-919-140B-9
US-07-919-140B-9
US-08-924-763-7
US-08-617-256-12
US-09-287-141-12
US-09-287-141-12
US-09-287-642-12
US-09-287-682-12
US-09-387-661-246
US-09-397-66-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ilarity 100.0%; Score 20; DB:
ilarity 100.0%; Pred. No. 0.2;
Conservative 0; Mismatches
                                                                                                                                                                                                                                              ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; OTHER INFORMATION: antisense sequence US-09-377-310-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 23, Application US/09377310B
; Patent No. 6133031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cogcoggeteacagtogteg 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 ccgcgggctcacagtggtcg
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Best Local Similarity
Lac 20; Conserve
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US-09-377-310-23
                                                                                                                                                                                                                                                                                             US-09-377-310-3
TYPE: DNA
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                                                                                ; Search time 417.38 Seconds (without alignments) 9.071 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                           to have a
being printed,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Description
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Sequence 5
Sequence 9
Sequence 1
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                                                                                                                                                                                                                                                                                                                                                                   /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/PcTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/PcTUS_COMB.seq:*
          GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-282-996-18
US-08-361-337-41
US-08-372-612-6
US-08-732-612-6
US-09-135-639-8
US-09-135-639-9
US-09-315-886C-5
US-09-315-886C-5
US-08-74-128-38
US-08-74-128-38
PCT-US95-05602-38
PCT-US95-05602-38
US-08-640-378-4
US-08-640-378-4
US-08-640-378-5
US-08-640-378-5
US-08-640-378-5
US-08-640-378-5
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PCT-US95-05816-34
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                                                                                                                                                                                                                     324599 segs, 94655562 residues
                                                                                  2001, 12:00:36
                                                                                                                                                                                                                                                                                                      Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                           · nucleic search, using sw model
                                                                                                                                                          ccgcgggctcacagtggtcg 20
                                                                                                                                                                                 IDENTITY_NUC Gapop 10.0 , Gapext 1.0
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                                                                                                                                US-09-757-100B-3
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Match Length
                                                                                                                                                                                                                                                                    length: 0
length: 50
                                                                                  October
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Maximum DB seq
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Sequence:
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APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF ETS-2 EXPRESSION
FILE REFERENCE: RTS-0063
CURRENT APPLICATION UNMBER: US/09/344,579
CURRENT FILING DATE: 1999-06-25
NUMBER OF SEQ ID NOS: 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 6, Application US/08732612;
Patent No. 5922583;
GENERAL INFORMATION:
APPLICANT: MORSEY, MOHAMAD A.
TITLE OF INVENTION: METHODS FOR PRODUCTION OF RECOMBINANT TITLE OF INVENTION: PLASMIDS
                                                                                                                                                                                                                                                                                                                                                                                                                 Score 13.2; DB 1;
Pred. No. 3.8e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 12.2; DB 3;
Pred. No. 1.1e+03;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: Antisense Oligonucleotide US-09-344-579-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: MORRISON & FOERSTER STREET: 2000 PENNSYLVANIA AVE., NW
                                                                                                                    REFERENCE/DOCKET NUMBER: 1414.634
TELECOMMUNICATION INFORMATION:
TELEPHONE: (404) 688-970
TELEFAX: (404) 688-980
INFORMATION FOR SEQ ID NO: 41:
SEQUENCE CHARACTERISTICS:
LENGTH: 33 base pairs
  APPLICATION NUMBER: US/08/361,337
                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear MOLECULE TYPE: Other nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 40, Application US/09344579; Patent No. 6054316; GENERAL INFORMATION:
                                                        ATTORNEY/AGENT INFORMATION:
NAME: Perryman, David G.
REGISTRATION NUMBER: 33,438
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82.4%;
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83.3%;
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Best Local Similarity 83.3'
Matches 15; Conservative
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                       nucleic acid
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Best Local Similarity
Matches 14; Conserv
                                           CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 5
US-09-344-579-40
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LENGTH: 18
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                                                                                                   75.0%; Score 15; DB 3; Length 15; 100.0%; Pred. No. 50; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Grentzmann, Guido
APPLICANT: Grentzmann, Guido
APPLICANT: Grentzmann, Raymond F.
APPLICANT: Atklins, John F.
TITLE OF INVENTION: Dual-Luciferase Reporter System
FILE REFERENCE: T5864.NP
CURRENT APPLICATION NUMBER: US/09/282,996
CURRENT FILING DATE: 1999-03-31
EARLIER APPLICATION NUMBER:
FARLIER FILING DATE:
NUMBER OF SEQ ID NOS: 18
SOFTWARE: WordPerfect 8.0
SEQ ID NO 18
: LENGTH: 31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Levenbook, Inessa S.
APPLICANT: Chumakov, Konstantin M.
APPLICANT: No. 5728519wood, Laurie P.
APPLICANT: Roninson, 1gor.
TITLE OF INVENTION: ASSAY FOR VIRULENT REVERTAN
TITLE OF INVENTION: ATTENUATED LIVE VACCINES
NUMBER OF SEQUENCES: 77
CORRESPONDENCE ADDRESS:
ADDRESSEE: NEEDLE & ROSENBERG, P.C.
STREET: 127 Peachtree Street, N.E.
        ; OTHER INFORMATION: antisense sequence US-09-377-310-23
                                                                                                                                                                                                                                                                                                                               Sequence 18, Application US/09282996 Patent No. 6143502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 41, Application US/08361337
Patent No. 5728519
GENERAL INFORMATION:
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US-09-282-996-18
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                                                                                                   Query Match 75.C
Best Local Similarity 100.
Matches 15; Conservative
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Best Local Similarity 88.2
Matches 15; Conservative
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COMPUTER READABLE FORM:
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STATE: Georgia
                                                                                                                                                                                                                                                                                                         -09-282-996-18/c
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US-08-361-337-41
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FEATURE:
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Gaps

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Indels

Length 18;

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Gaps

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Indels

Length 33;

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APPLICANT: LUNNEN, MICHEL A.,
APPLICANT: DALTON MICHEL A.,
APPLICANT: DALTON MICHEL A.,
APPLICANT: WILSON, GEOFFREY G.
APPLICANT: XU, SHUANG-YONG
TITLE OF INVENTION: Method For Cloning And Producing The Avai Restriction
TITLE OF INVENTION: Recombinant Avai Restriction Endonuclease
FILE REFERENCE: Avai
CURRENT APPLICATION NUMBER: US/09/135,639
CURRENT FILING DATE: 1998-08-18
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 10, Application US/09135639

Sequence 10, Application US/09135639

Patent No. 6004793

GENERAL INFORMATION:
APPLICANT: LONNEN: LONNEN:
APPLICANT: DALFON MICHAEL A.,
APPLICANT: MILSON, GEOFFREY G.
APPLICANT: XU, SHUANG-YONG
TITLE OF INVENTION: Method For Cloning And Producing The Aval Restriction
TITLE OF INVENTION: Endonuclease In E. coli And Purification Of The
TITLE OF INVENTION: Recombinant Aval Restriction Endonuclease
FILE REFERENCE: Aval
                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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Pred. No. 1.2e+03;
0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                Score 12.2; DB 2;
Pred. No. 1.2e+03;
0; Mismatches 3;
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IELEFAX: (202) 887-0763
TELEX: 90-4030 MRSNFOERSWSH
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 39 base pairs
TEPE: nucleic acid
STRANDEDNESS: single
FOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 8, Application US/09135639; Patent No. 6004793
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; ORGANISM: Anabaena variabilis
US-09-135-639-8
                                                                                                                                                                                                                                  prim_transcript
                                                                                                                                                                                                                                                                                                                                                                                  61.0%;
82.4%;
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Best Local Similarity 82.4
Matches 14; Conservative
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CTHER INFORMATION:
COTHER INFORMATION:
US-08-732-612-11
                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 14; Conserv
                                                                                                                                                                                                                              NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 9
US-09-135-639-10
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Patent No. 522283
GENERAL INFORMATION:
APPLICANT: MORSEY, MOHAMAD A.
TITLE OF INVENTION: METHODS FOR PRODUCTION OF RECOMBINANT
TITLE OF INVENTION: PLASMIDS
NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
STREET: 2000 PENNSYLVANIA AVE., NW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                          COMPUTER RELDABLE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/732,612
FILING DATE: 16-OCT-1996
SCHASSIFICATION: 435-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
ZIP: 20006-1812
COMPUTER READABLE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/732,612
FILING DATE: 16-OCT-1996
CLASSIFICATION: 435
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Pred. No. 1.2e+03;
0; Mismatches 3;
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                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: MURASHIGE, KATE H.
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 29311-20006.21
TELECOMMUNICATION INFORMATION:
TELEPAX: (202) 887-0763
TELEFAX: (202) 887-0763
IELEFAX: 90-4030 MRSNFOERSWSH
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
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NAME: MURASHIGE, KATE H.
REGISTRATION NUMBER: 29,959
REFERENCET NUMBER: 29311-20006.21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 887-1500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: prim_transcript
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Best Local Similarity 82.4%;
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 31 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LOCATION: 1.31
OTHER INFORMATION:
OTHER INFORMATION:
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SOFTWARE: PatentIn Release #1.0, Version #1.30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 13, Application US/08640378; Patent No. 5811269; GENERAL INFORMATION:
                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Highlander, Steven L
REGISTRATION NUMBER: 37,642
REFERENCE/DOCKET NUMBER: ARSB
TELECOMMUNICATION INFORMATION:
TELEFHONE: 512/418-3000
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
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NAME: Fugit, Donna R.
REGISTRATION NIMBER: 32,135
REFERENCE/DOCKET NUMBER: P-.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 ccgcgggctcacagtggtcg 20
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                       CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                            LENGTH: 41 base pairs
TYPE: nucleic acid
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TYPE: nucleic acid
STRANDEDNESS: single
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CLASSIFICATION: 435
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Matches 15; Conserv
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                                                                                                                                                                    Length 40;
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APPLICANT: Khvorova, Anastasia
APPLICANT: Yarus, Michael
TITLE OF INVENTION: RNA Channels in Biological Membranes
FILE REFERENCE: UTC-03444
CURRENT APPLICATION NUMBER: US/09/315,886C
CURRENT FILING DATE: 1999-05-20
PRIOR APPLICATION NUMBER: 60/086,492
PRIOR FILING DATE: 1998-05-22
NUMBER OF SEQ ID NOS: 32
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 5
LENGTH: 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    61.0%; Score 12.2; DB 4; Length 50; 64.7%; Pred. No. 1.2e+03;
Live 3; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Michael, Mancy M
APPLICANT: Accavitt, Marianne
APPLICANT: Thompson, Craig B
TITLE OF INVENTION: MONOCLONAL ANTIBODIES
NUMBER OF SEQUENCES:
ADDRESSE: Annold, White & Durkee
                                                                                                                                                                                                          3; Indels
                                                                                                                                                                    Score 12.2; DB 3;
Pred. No. 1.2e+03;
); Mismatches 3;
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                        Sequence 5, Application US/09315886C
Patent No. 6225063
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Patent No. 6143559
GENERAL INFORMATION:
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                                                                                            ; ORGANISM: Anabaena variabilis
US-09-135-639-10
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Best Local Similarity 82.4%;
Matches 14; Conservative
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NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 10
TENGTH: 40
TYPE: DNA
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Best Local Similarity
Matches 11; Conserve
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STATE: Texas
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US-08-751-359-9/c
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US-09-315-886C-5
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Gaps
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APPLICANT: Schram, James L.
APPLICANT: Schram, James L.
APPLICANT: Howard, Deborah R.
APPLICANT: Wargaret S.
APPLICANT: Wright, David J.
TITLE OF INVENTION: DETECTION OF MYCOBACTERIA BY MULTIPLEX
TITLE OF INVENTION: UUCLEIC ACID AMPLIFICATION
CORRESPONDENCE: 27
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Richard J. Rodrick, Becton Dickinson and ADDRESSEE: Company STREET: 1 Becton Drive CITY: Franklin Lakes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/640,378
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           . 1.5e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     60.0%; Score 12; DB 3; 75.0%; Pred. No. 1.5e+03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
APPLICATION NUMBER: US/08/751,359
FILING DATE: Concurrently Herewith
CLASSIFICATION: 510
                                                                                                                                                         ARSB: 504
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APPLICANT: Solomon, N. T.

APPLICANT: Solomon, N. T.

TITLE OF INVENTION: Materials and Methods for the Detection of TITLE OF INVENTION: Mycobacteria tuberculosis
NUMBER OF SEQUENCES: 76

CORRESPONDENCE ADDRESS:
ADDRESSEE: Abbott Laboratories
STREET: One Abbott Laboratories
STREET: One Abbott Park Road
CITY: Abbott Park Road
CITY: Abbott Park Road
CITY: Abbott Park Road
CITY: Abbott Park Road
CONPUTER: Illinois
COMPUTER: EMP PC compatible
OMPUTER: IBM PC compatible
OMPUTER: Jame PC compatible
OMPUTER: IBM PC compatible
OMPUTER: 
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NUMBER OF SEQUENCES: 76
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 11.8; DB 1; Length 22;
Pred. No. 1.8e+03;
0; Mismatches 2; Indels
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STREET: One Abbott Park Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Leckie, G.W.
APPLICANT: Davis A.H.
APPLICANT: Semple-Facey, I.E.
APPLICANT: Manlove, M.T.
APPLICANT: Solomon, N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; TOPOLOGY: linear
; MOLECULE TYPE: Synthetic DNA
US-08-774-128-38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          59.0%;
86.7%;
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 22 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
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Patent No. 5631130
GENERAL INFORMATION
GENERAL INFORMATION
APPLICANT: Leckie, G. W.
APPLICANT: Semple-Facey, I. E.
APPLICANT: Solomon, N. A.
TITLE OF INVENTION: Materials and Methods for the Detection of
TITLE OF INVENTION: Mycobacteria tuberculosis
NUMBER OF SEQUENCES: 76
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Score 11.8; DB 1; Length 16;
Pred. No. 1.8e+03;
0; Mismatches 2; Indels
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/242,403A
FILING DATE: May 13, 1994
CLASSIFTCATION: 435
ATTORNEY,AGENT INFORMATION:
NAME: Thomas D. Brainard
REGISTRATION NUMBER: 32,459
REFRERENCE/DOCKET NUMBER: 5370.US.01
TELECHONE: 708/937-4884
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                %308-774-128-38
Sequence 38, Application US/08774128
Patent No. 5786149
GENERAL INFORMATION:
APPLICANT: Leckie, G. W.
APPLICANT: Davis, A. H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Abbott Laboratories
STREET: One Abbott Park Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible OPERATING SYSTEM: PCDOS/MSDOS
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  59.0%;
86.7%;
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SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
LYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Floppy disk
                             Best_Local Similarity 86.7
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Abbott Park
STATE: Illinois
COUNTRY: USA
ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy di
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Best Local Similarity 86.7
Matches 13; Conservative
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TELEFAX: 708/938-2623
                                                                                                                  1 ccgcgggctcacagt 15
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                                                                                                                                               15 CCGCATGCTCACAGT 1
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ADDRESSEE: Abbott La
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
                                                                                                                                                                                                                                                                             US-08-242-403A-38
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     Query Match
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Gaps

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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC_DOS/MS_DOS
SOCTWARE: WOTOPErfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/05602
FILING DATE: May 13, 1994
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAMME: Thomas D. Brainard
FEGISTRATION NUMBER: 5370.PC.01
FELECOMMUNICATION INFORMATION:
TELEPHONE: 708/937-4884
TELEPHONE: 708/938-2623
TELECOMMUNICATION SY 38:
TELEPHONE: TOR/938-2623
TELEPHONE: TOR/938-263
INFORMATION FOR SEQ ID NO: 38:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
TOPOLOGY: linear
MOLECULE TYPE: Synthetic DNA
PCT-US95-05602-38
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Gaps

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Query Match

59.0%; Score 11.8; DB 5; Length 22;
Best Local Similarity 86.7%; Pred. No. 1.8e+03;
Matches 13; Conservative 0; Mismatches 2; Indels

1 ccgcgggctcacagt 15 |||| ||||||||||| 7 CCGCACGCTCACAGT 21

9 9 Search completed: October 2, 2001, 16:03:44 Job time: 14588 sec

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/cgn2_6/ptodata/1/pna/US6010_COMB.seq:*
/cgn2_6/ptodata/1/pna/US6011_COMB.seq:*
                GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 1.0
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20
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Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description	Sequence 11, Appl Sequence 11, Appl	Ξ	m	9034,	32952	equence 32964	Sequence 35	Ξ,	7	31, App	7	~;		0/075	Sequence 439150,		Sequence 329525,		Sequence 22343, A		Sequence 51615, A				57516	15388	Sequence	Ĺ	71111,	63810,	Sequence 63910, A	4.			٦,	95260	20118	3204	335	3972	eguence 125	40	Sequence 215290,
SUMMARIES	ID	US00-18999-1 09-377-310-1	7-100B-11	-60-233-166-320	50-233-357-9034	50-233-166-32952	-16	33-166-32964	JS00-18999-3	9-377-310-3	09-757-100B-31	00-234-017-2	00-233-620-	US-60-234-01/-142091	20-234-01/	50-234-01/-	-01-532-00 -033-100-	US-60-233-166-329323	20-780-08	50-234-017-	19-660-220-546	233-166	50-233-166-4098	S-60-234-017-2418	S-60-234-017-4288	8-60-234-017-57516	8-60-234-017-1538	5-60-234-017-52354	29-7	US-09-660-220-71111	166-	US-60-233-166-63910	077	05-08-660-7711755	- 770-	S-60-232-638-	S-60-233-166-95260	5-60-233-166-20118	S-60-233-166-320	S-60-233-166-33544	8-60-233-166-39724	-60-233-620-12535	-60-234-017-17407	US-60-234-017-215290
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GENERAL INFORMATION:
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GENERAL INFORMATION:

APPLICANT: Monia, Brett P.

APPLICANT: Garade, William A.

TITLE OF INVENTION: Expression

TITLE OF INVENTION: Expression

TITLE OF INVENTION: Expression

TITLE OF INVENTION: Expression

CURRENT APPLICATION UNDMER: US/09/377,310A

CURRENT PILING DATE: 1999-08-19

NUMBER OF SEQ ID NOS: 43

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 11

LENGTH: 20
                                                                                                                                                    APPLICANT: Moria, Brett P.
APPLICANT: Gaarde, William A.
APPLICANT: Nero, Pamela S.
TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase TITLE OF INVENTION: Expression
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                ALIGNMENTS
                                                                                                                                                                                                                                                FILE REFERENCE: ISPH-0476
CURRENT APPLICATION NUMBER: PCT/US00/18999
CURRENT FILING DATE: 2000-07-13
PRIOR APPLICATION NUMBER: 09/377,310
PRIOR FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SOFTWARE: PATENTIN VET. 2.0
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Pred. No.
                                                                                                                                             Isis Pharmaceuticals, Inc.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: antisense sequence PCT-US00-18999-11
                                                                                                  Sequence 11, Application PC/TUS0018999 GENERAL INFORMATION:
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US-09-757-100B-11
; Sequence 11, Application US/09757100B
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ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 20; Conservative
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Best Local Similarity 100.
Matches 20; Conservative
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LENGTH: 20
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                                                        TITLE OF INVENTION: Antisense Modulation of Focal Adhesion Kinase TITLE OF INVENTION: Expression FILE REFERENCE: ISPH-0533
FILE REFERENCE: ISPH-0533
CURRENT APPLICATION NUMBER: US/09/757,100B
CURRENT FILING DATE: 2001-03-15
PRIOR PILING DATE: 1999-08-19
PRIOR FILING DATE: 1999-08-19
PRIOR FILING DATE: 2000-07-13
NUMBER OF SEQ ID NOS: 44
SOFTWARE: PATENTIN VET: 2.0
SEQ ID NO 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-60-233-166-320910/C
) Sequence 3.20910, Application US/60233166
) GENERAL INFORMATION
) APPLICANT: Mittmann
) APPLICANT: Affymetrix, Inc.
) TITLE OF INVENTION: Methods of Genetic Analysis of the Rat TITLE OF INVENTION: Genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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APPLICANT: Affymetrix, Inc.
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of Rat
TITLE OF INVENTION: Neurobiology
FILE REFERENCE: 3114
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Pred. No. 5.2e+02;
0; Mismatches 2;
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Pred. No. 2.9;
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CURRENT APPLICATION NUMBER: US/60/233,166
CURRENT FILING DATE: 2000-10-24
NUMBER OF SEO ID NOS: 420907
SOFTWARE: FastSEQ for Windows Version 4.0
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; DATABASE ACCESSION NUMBER: GenBank L08493
US-60-233-166-320910
                                                                                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION; antisense sequence US-09-757-100B-11
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GENERAL INFORMATION:
APPLICANT: Mittmann
APPLICANT: Affymetrix, Inc.
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APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
APPLICANT: Nero, Pamela S.
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Matches 17; Conservative
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Best Local Similarity 100.
Matches 20; Conservative
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US-60-233-357-9034/c
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LENGTH: 25
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                                           Length 25;
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Sequence 329644, Application US/60233166
GENERAL INFORMATION:
APPLICANT: Mittmann
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of the Rat
FILLE REFERENCE: 3112
CURRENT APPLICATION UNMBER: US/60/233,166
CURRENT FILING DATE: 2000-10-24
NUMBER OF SEQ ID NOS: 420907
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          75.0%; Score 15; DB 1; Length 15; 100.0%; Pred. No. 1.3e+03; Live 0; Mismatches 0; Indels
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                                           Query Match 76.0%; Score 15.2; DB 55; Best Local Similarity 85.0%; Pred. No. 1.1e+03; Matches 17; Conservative 0; Mismatches 3;
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76.0%; Score 15.2; DB 55;
Best Local Similarity 85.0%; Pred. No. 1.1e+03;
Matches 17; Conservative 0; Mismatches 3;
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; PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank M31725
US-60-233-166-329644
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 31, Application PC/TUS0018999 ; GENERAL INFORMATION:
                                                                                                                                 1 agtacccaggtgagtcttag 20
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Best Local Similarity 100.
Matches 15; Conservative
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PCT-US00-18999-31
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APPLICANT:
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APPLICANT:
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LENGTH: 15
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US-60-233-166-329526
; Sequence 329526
; Sequence 329526
; GENERAL INFORMATION:
; APPLICANT: Mittmann
; APPLICANT: Mittmann
; TILLE OF INVENTION: Methods of Genetic Analysis of the Rat
; TILLE OF INVENTION: Genome
; TILLE REFERENCE: 3112
; CURRENT FALLIA DATE: 2000-10-24
; NUMBER OF SEQ ID NOS: 420907
; SOFTWARE: FastSEQ for Windows Version 4.0
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GENERAL INFORMATION:
APPLICANT: Mithmann
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Genome
FILE REFERENCE: 3112
CURRENT PEPLICATION NUMBER: US/60/233,166
CURRENT FILING DATE: 2000-10-24
NUMBER OF SEQ ID NOS: 420907
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 329643
LENGTH: 25
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Pred. No. 5.2e+02;
0; Mismatches 2;
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                CURRENT FILING DATE: 2000-10-24
NUMBER OF SEQ ID NOS: 21305
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 9034
LENGTH: 25
                                                                                                                                                                    ; PUBLICATION INFORMATION:
; DATABASE ACCESSION NUMBER: Genbank L08493
US-60-233-357-9034
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; DATABASE ACCESSION NUMBER: Genbank M31725
US-60-233-166-329526
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US-60-233-166-329643
CURRENT APPLICATION NUMBER: US/60/233,357
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Best Local Similarity 89.5%;
Matches 17; Conservative (
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|| || || || || || || || || || 3 agaacccaggtaaatcttag 22
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                                                                                                                          TYPE: DNA
ORGANISM: Rattus norvegicus
                                                                                                                                                                                                                                                                                                                                                                                                       24 GTAATCAGGTGAGTCTTAG 6
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LENGTH: 25
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Sequence 22548, Application US/60234017
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88.9%;
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5 gtaaccatgtgagtctta 22
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Best Local Similarity 88.9
Matches 16; Conservative
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Best Local Similarity 93.8
Matches 15; Conservative
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10 cccagctgagtcttag 25
                                                                                                                                                                                                                                   TYPE: DNA ORGANISM: Mus musculus
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                    GENERAL INFORMATION:
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US-60-233-620-18028
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GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
APPLICANT: Garde, William A.
APPLICANT: Nero, Pamela S.
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0533
CURRENT APPLICATION WUMBER: US/09/757,100B
CURRENT FILING DATE: 2001-03-15
PRIOR FILING DATE: 1999-08-19
PRIOR PILING DATE: 2000-07-13
NUMBER OF SEQ ID NOS: 44
SOFTWARE PATENTION NUMBER: PCT/US00/18999
NUMBER OF SEQ ID NOS: 44
SOFTWARE PATENTIN UNIVER: 2000-07-13
NUMBER OF SEQ ID NOS: 44
SEQ ID NO 31
LENGTH: 15
                                                                                                                                    APPLICANT: Monia, Brett P.
APPLICANT: Monia, Brett P.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: Antibense Modulation of Focal Adhesion Kinase
TITLE OF INVENTION: Expression
FILE REFERENCE: ISPH-0389
CURRENT FILING DATE: 1999-08-19
NUMBER OF SEQ ID NOS: 43
SECTRARE: Patentin Ver. 2.0
SEQ ID NO 31
LENGTH: 15
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7. 1.3e+03;
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100.0%; Pred. No. 1.3e+03;
Live 0; Mismatches 0;
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                                                                                 US-09-377-310-31; Sequence 31, Application US/09377310A; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 15; Conservative
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Matches 15; Conservative
3 tacccaggtgagtct 17
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US-60-234-017-22548
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APPLICANT: Mittmann, M
APPLICANT: Affymetrix, Inc.
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of Mus;
TITLE OF INVENTION: musculus
FILE REFERENCE: 3115
CURRENT FILE ADDATE: 1000-09-20
NUMBER OF SEQ ID NOS: 605887
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2248
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GENERAL INFORMATION:
APPLICANT: Mittmann, M
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of Mus
TITLE OF INVENTION: musculus
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GENERAL INFORMATION:
APPLICANT: Mittmann
APPLICANT: Affametrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis of
TITLE OF INVENTION: Arabidopsis thaliana
FILE REFERENCE: 3116
CURRENT APPLICATION NUMBER: US/60/233,620
CURRENT FILING DATE: 2000-10-24
                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 14.8; DB 55;
Pred. No. 1.8e+03;
0; Mismatches 2;
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Pred. No. 2.9e+03;
0; Mismatches 1;
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US-60-234-017-22548
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; DATABASE ACCESSION NUMBER: Genbank AC005724
US-60-233-620-18028
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 18028
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CURRENT FILING DATE: 2000-09-20
NUMBER OF SEQ ID NOS: 605887
SOFTWARE: FastSEQ for Windows Version 4.0
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71.0%; Score 14.2; DB 55; Length 25;
Best Local Similarity 84.2%; Pred. No. 3.7e+03;
Matches 16; Conservative 0; Mismatches 3; Indels (
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71.0%; Score 14.2; DB 55; Length 25;
Best Local Similarity 84.2%; Pred. No. 3.7e+03;
Matches 16; Conservative 0; Mismatches 3; Indels (
                                                                                                                                                                                                                                                                                                                                                       RESULT 15
US-60-234-017-420702
Sequence 420702. Application US/60234017
GENERAL INFORMATION:
APPLICANT: Mittmann, M
TITLE OF INVENTION: Methods of Genetic Analysis of Mus
TITLE OF INVENTION: Methods of Genetic Analysis of Mus
TITLE OF INVENTION: Musculus
FILE REFERENCE: 3115
CURRENT FILING DATE: 2000-09-20
NUMBER OF SEQ ID NOS: 605887
SOTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 420702
LENGTH: 25
                 LENGTH: 25
TYPE: DNA
ORGANISM: MUS musculus
PUBLICATION INFORMATION:
1 DATABASE ACCESSION NUMBER: Genbank AI591702
US-60-234-017-142091
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PUBLICATION INFORMATION:
CATABASE ACCESSION NUMBER: Genbank AU024707
US-60-234-017-420702
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1 agtcccaagtgtgtctta 19
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; LENGTH: 25
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Search completed: October 2, 2001, 21:50:14 Job time: 24527 sec

12604, A 12605, A 12606, A 12609, A 12609, A 12610, A 12611, A 12613, A

49, Appl 2830, Ap 1301, Ap 308, App 7, Appli 4142, Ap 4149, Ap

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Run on:

Searched:

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Sequence 65, Application US/09857581
GENERAL INFORMATION:
APPLICANT: E. I. du Pont de Nemours and Company
ITILE OF INVENTION: Nucleic Acid Sequences Encoding Isoflavone Synthase
FILE REPRENCE: BB1339 PCT
CURRENT ELLING DATE: 2001-06-05
CURRENT FILING DATE: 1999-01-27
PRIOR APPLICATION NUMBER: 60/117,769
PRIOR FILING DATE: 1999-01-27
PRIOR PRIOR APPLICATION NUMBER: 60/144,783
PRIOR PLING DATE: 1999-01-27
PRIOR FILING DATE: 1999-00-24
NUMBER OF SEQ ID NOS: 66
SOFTWARE: Microsoft Office 97
SEQ ID NO 65
LENGTH: 25
                                                                       Sequence Seq
                                                                                                                                                                                                                                                                                                                                     Sequence Sequence
                                                                                                                                                                                                                                                                                                                                                                                                            Sequence S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 122, Application US/09672217
GENERAL INFORMATION:
APPLICANT: FELL, JACK
APPLICANT: DIAZ, MARA
TILLE OF INVENTION: METHOD OF IDENTIFYING PATHOGENIC CRYPTOCOCCI
FILLE REPRENCE: 0862220143
CURRENT APPLICATION NUMBER: US/09/672,217
CURRENT APPLICATION NUMBER: 60/156,598
PRIOR PILLING DATE: 1999-09-29
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                   US-09-648-103B-51

US-09-666-108-12604

US-09-866-108-12605

US-09-866-108-12605

US-09-866-108-12608

US-09-866-108-12609

US-09-866-108-12610

US-09-866-108-12611

US-09-866-108-12611

US-09-866-108-12611

US-09-866-108-12611

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US-09-866-108-12611

US-09-866-108-13611

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87.5%; Pred. No. 2.1e+03;
ive 0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Artificial Sequence
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4; Conservative
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Best Local Simi
Matches 14;
FEATURE:
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Sequence 122, App
Sequence 223, App
Sequence 16, Appl
Sequence 4143, Ap
Sequence 4144, Ap
Sequence 4146, Ap
Sequence 4146, Ap
Sequence 4146, Ap
Sequence 4141, Ap
Sequence 42408, Ap
Sequence 69, Appl
Sequence 69, Appl
Sequence 69, Appl
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Sequence 7713, Ap
Sequence 18, Appl
Sequence 2232, Ap
Sequence 2323, Ap
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Sequence 115, App
Sequence 115, App
Sequence 82, Appl
                                                                                                                                                                                      (without alignments)
19.290 Million cell updates/sec
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                                                                                                                                                                Search time 2173.58 Seconds
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1: /cgn2_6/ptodata/2/pna/PCT_NEW_COMB.seq:*

2: /cgn2_6/ptodata/2/pna/USOG_NEW_COMB.seq:*

3: /cgn2_6/ptodata/2/pna/USOR_NEW_COMB.seq:*

5: /cgn2_6/ptodata/2/pna/USOR_NEW_COMB.seq:*

6: /cgn2_6/ptodata/2/pna/USOR_NEW_COMB.seq:*

7: /cgn2_6/ptodata/2/pna/USOR_NEW_COMB.seq:*

8: /cgn2_6/ptodata/2/pna/USOR_NEW_COMB.seq:*

8: /cgn2_6/ptodata/2/pna/USOR_NEW_COMB.seq:*
                      GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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US-09-672-217-122
US-09-6672-217-223
US-09-866-108-4144
US-09-866-108-4144
US-09-866-108-4145
US-09-866-108-4145
US-09-866-108-4146
US-09-866-108-4148
US-09-866-108-4148
US-09-866-108-4148
US-09-866-108-4181
US-09-252-457-42408
US-09-868-806-69
US-09-801-274-1705
US-09-866-108-7712
US-09-866-108-7712
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US-09-895-007-115
US-09-920-313-115
US-09-659-845A-82
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Listing first 45 summaries

    nucleic search, using sw model

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Match Length
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Sequence:
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Gaps

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Result Ş.

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APPLICANT: RANK, David R.

APPLICANT: HANKEL, David R.

APPLICANT: SHANKON, MARK

TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
CURRENT APPLICATION WINDER: US/09/866,108

CURRENT FILING DATE: 2001-05-25

PRIOR PILING DATE: 2000-05-26

PRIOR APPLICATION WINDER: US 60/207,456

PRIOR FILING DATE: 2000-10-04

PRIOR FILING DATE: 2000-10-04

PRIOR FILING DATE: 2000-10-04

PRIOR FILING DATE: 2001-01-30

PRIOR PLING DATE: 2001-01-30

PRIOR PLING DATE: 2001-01-30

PRIOR PLING DATE: 2001-01-30

PRIOR PRILING DATE: 2001-01-30

PRIOR PLING DATE: 2001-01-30

PRIOR PRILING DATE: 2001-01-30

PRIOR PLING DATE: 2000-09-21

PRIOR PLING DATE: 2001-01-30

PRIOR PRILICATION NUMBER: US 60/234,687

PRIOR PRIOR PLING DATE: 2001-01-30

PRIOR PRILICATION NUMBER: US 60/226,860

PRIOR PRILICATION NUMBER: US 60/226,860

PRIOR PRILICATION NUMBER: US 60/226,860
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                            Length 22;
                              ; Db -,
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Pred. No. 5.9e+03;
0; Mismatches 5;
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                            Score 12;
Pred. No. 5
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SEQ ID NO 4143
LENGTH: 25
                                                                                                                                                                                                                                                                                                 Sequence 4143, Application US/09866108; GENERAL INFORMATION:
APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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75.0%;
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75.08;
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                                                   Best Local Similarity 75.0
Matches 15; Conservative
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CORGANISM: Homo sapiens
US-09-866-108-4143
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Best Local Similarity
                            Query Match
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TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES ENCODED THEREBY DISTANTLY HOMOLG
TITLE OF INVENTION: HEPARANASE
FILE REFERENCE: 00/20105
CURRENT FILIA DATE: 2000-01-07
PRIOR APPLICATION NUMBER: 60/140,801
PRIOR APPLICATION NUMBER: 60/140,801
PRIOR FILING DATE: 1999-06-25
NUMBER OF SEQ ID NOS: 25
SOFTWARE: PATHOLICATION 3.0
SEQ ID NO 16
LENGTH: 22
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Sequence 223, Application US/09672217

GENERAL INFORMATION:

APPLICANT: FELL, JACK

APPLICANT: BIAZ, MARA

APPLICANT: MCCABE, MEADE

TILLE OF INVENTION: METHOD OF IDENTIFYING PATHOGENIC CRYPTOCOCCI

FILE REPERENCE: 086222/013

CURRENT FILING DATE: 2000-09-29

PRIOR PILING DATE: 1999-09-29

PRIOR FILING DATE: 1999-09-29

NUMBER: OF EXQ ID NOS: 301

SOFTWARE: Patentin Ver: 2.1

SEQ ID NO 223

LENGTH: 33
                                                                                                                                                                         ; OTHER INFORMATION: Description of Artificial Sequence: Primer/Probe US-09-672-217-122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: Description of Artificial Sequence: Primer/Probe US-09-672-217-223
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TC-1050-00358-16

Sequence 16, Application PC/TUS0000358
; GENERAL INFORMATION:
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; ORGANISM: Synthetic Oligonucleotide
PCT-USOO-00358-16
                                                                                                   TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
NUMBER OF SEQ ID NOS: 301
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 122
LENGTH: 33
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Best Local Similarity 92.9
Matches 13; Conservative
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Best Local Similarity 92.9
Matches 13; Conservative
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APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
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              THE REPERENCE: ADOMICA-7

CURRENT APPLICATION NUMBER: US/09/866,108

CURRENT APPLICATION NUMBER: US/09/866,108

CURRENT APPLICATION NUMBER: US/09/866,108

PRIOR PILING DATE: 2000-05-26

PRIOR FILING DATE: 2000-06-26

PRIOR FILING DATE: 2000-09-27

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR APPLICATION NUMBER: PCT/US01/00666

PRIOR PILING DATE: 2001-01-30

PRIOR FILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/US01/00669

PRIOR FILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/US01/00661

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30
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75.0%; Pred. No. 5.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Aeomica Sequence Listing Engine
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APPLICATION NUMBER: US 60/236,359
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 4146, Application US/09866108 GENERAL INFORMATION:
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PENN, Sharron G.
HANZEL, David K.
RANK, David R.
CHEN, Wensheng
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Best Local Similarity 75.0
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-866-108-4146
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APPLICANT:
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                                                                                                                                                                                                                       APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, MATK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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ches 5;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: GB 24263.6
PRIOR FLILING DATE: 2000-10-04
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR PELIANG NUMBER: PCT/US01/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00667
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APPLICATION NUMBER: PCT/US01/00663
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APPLICATION NUMBER: PCT/US01/00662
FILING DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00664
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FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
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                    Sequence 4144, Application US/09866108 GENERAL INFORMATION:
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APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
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; ORGANISM: Homo sapiens
US-09-866-108-4144
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Best Local Similarity
Matches 15; Conserv
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APPLICANT: GHEN, WALESLEURY
APPLICANT: GHEN, WALESLEURY
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEGNICA-7
CURRENT APPLICATION NUMBER: US/09/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-00-04
PRIOR FILING DATE: 2000-00-04
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2001-01-30
PRIOR PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR PRIOR APPLICATION NUMBER: PCT/US01/00669
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Pred. No. 5.9e+03;
Transanhes 5; Indels
                            FRIOR AFFLICATION NUMBER: 2001-01-30
PRIOR FILING DATE: 2001-01-30
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PRIOR FILING DATE: 2001-01-30
PRIOR PILING DATE: 2001-01-30
PRIOR FILING DATE: 2001-01-30
PRIOR PILING DATE: 2000-01-31
PRIOR PILING DATE: 2000-09-21
PRIOR PILING DATE: 2000-09-21
PRIOR FILING DATE: 2000-09-21
PRIOR FILING DATE: 2000-09-21
PRIOR FILING DATE: 2000-09-31
PRIOR FILING DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Acomica Sequence Listing Engine
SEQ ID NO 4447
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            APPLICATION NUMBER: PCT/US01/00668
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PRIOR APPLICATION NUMBER: PCT/US01/00665
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PRIOR APPLICATION NUMBER: PCT/US01/00668
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APPLICATION NUMBER: PCT/US01/00663
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75.0%;
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Best Local Similarity 75.0°
Matches 15; Conservative
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; ORGANISM: Homo sapiens
US-09-866-108-4147
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APPLICANT: 31, Yonggang
APPLICANT: GU, Yizhong
APPLICANT: GU, Yizhong
APPLICANT: HANZEL, David K.
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REPERENCE: ADDILCA-7
CURRENT APPLICATION NUMBER: US/99/866,108
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00666
PRIOR PILING DATE: 2001-01-30
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PRIOR APPLICATION NUMBER: PCT/USO1/00666
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/USO1/00667
PRIOR APPLICATION NUMBER: PCT/USO1/00667
PRIOR PILING DATE: 2001-01-30
PRIOR PELLING DATE: 2001-01-30
PRIOR PELLING DATE: 2001-01-30
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PRIOR PILING DATE: 2001-01-30
PRIOR PELLING DATE: 2001-01-30
PRIOR PILING DATE: 2000-09-21
PRIOR PILING DATE: 2000-09-21
PRIOR PILING DATE: 2000-09-21
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SOFTWARE: Aeomica Sequence Listing Engine
SEQ ID NO 4146
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00669
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APPLICATION NUMBER: PCT/US01/00665
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75.0%;
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Best Local Similarity 75.0
Matches 15; Conservative
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US-09-866-108-4146
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APPLICANT: Watson, James D
APPLICANT: Watson, James G
TITLE OF INVENTION: Polynucleotides, polypeptides expressed
TITLE OF INVENTION: by the polynucleotides and methods for their use.
FILE REPERBNE: 1050U2
CURRENT APPLICATION NUMBER: US/09/724,671
CURRENT FILING DATE: 2000-11-28
PRIOR APPLICATION NUMBER: US. No. 60/171,678
PRIOR FILING DATE: 1999-12-23
NUMBER OF SEC ID NOS: 21907
SOFTWARE: PESTSEQ for Windows Version 4.0
SEQ ID NO 10607
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Sequence 42408, Application US/60253457

Sequence 42408, Application US/60253457

GENERAL INFORMATION:
TITLE OF INVENTION: Polynuclectides, isolated from plants;
TITLE OF INVENTION: and methods for their use.
FILE REFERENCE: 1054P2

CURRENT APPLICATION NUMBER: US/60/253,457

CURRENT FILIG DATE: 2000-11-27

NUMBER OF SEC ID NOS: 48893

SOFTWARE: FastSEQ for Windows Version 4.0

FEMALET OF THE PROBLEM OF 
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                                                                                                       Length 17;
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Pred. No. 1.1e+04;
0; Mismatches 4;
                                                                                                       Score 11.8; DB 6;
Pred. No. 7.1e+03;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                     RESULT 13
Second 13
Sequence 10607, Application US/09724671
GENERAL INFORMATION:
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77.8%;
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22 AGTGTCCAGGTCAGTGTT 5
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Best Local Similarity 77.8
Matches 14; Conservative
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Best Local Similarity 77.8
Matches 14; Conservative
                                                                                                                                                                    Conservative
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CORGANISM: Pinus radiata
US-60-253-457-42408
                                                                                                       Query Match
Best Local Similarity
Matches 13; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
CRGANISM: Mouse
US-09-724-671-10607
                US-09-546-745A-4802
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APPLICANT: Zwick, Michael
APPLICANT: Zwick, Michael
APPLICANT: Paveco, Pan
APPLICANT: McSwiggen, Jim
TITLE OF INVENTION: Regulation of Repressor Genes using Nucleic Acid Molecules
FILE REFERENCE: 237/131
CURRENT APPLICATION NUMBER: US/09/546,745A
CURRENT FILING DATE: 2000-04-11
NUMBER OF SEQ ID NOS: 7043
SOFTWARE: PatentIn version 3.0
SEQ ID NO 40802
LENGTH: 17
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6.4e+03;
5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                         60.0%; Score 12; DB 6; Length 25; 75.0%; Pred. No. 5.9e+03;
.ive 0; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-60-252-833-24181/C
| Sequence 24481, Application US/60252833
| GENERAL INFORMATION:
| APPLICANT: Glenn, Matthew | APPLICANT: Glenn, Matthew | APPLICANT: Grigor, Murray R
| TITLE OF INVENTION: Compositions isolated from bovine | TITLE OF INVENTION: Compositions do nethods for their use: | FILE REFERENCE: 1652P2 | CURRENT APPLICATION NUMBER: US/60/252,833 | CURRENT FILING DATE: 2000-11-21 | NUMBER OF SED ID NOS: 43535 | SEQ ID NO 24181 | LENGTH: 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  60.0%; Score 12; DB ilarity 75.0%; Pred. No. 6.46 Conservative 0; Mismatches
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
PRIOR PELICATION NUMBER: US 60/266,860
PROR PILING, DATE: 2001-02-05
NUMBER OF SEQ ID NOS: 15752
SOFTWARE: Aeomica Sequence Listing Engine
SEQ ID NO 4148
LENGTH: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 4802, Application US/09546745A GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 agtacccaggtgagtcttag 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21 AGTACTCAGGGAAGCTTTAG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 60.0°
Best Local Similarity 75.0°
Matches 15; Conservative
                                                                                                                                                                                                                                                                                     TYPE: DNA
CORGANISM: Homo sapiens
US-09-866-108-4148
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ouery Match
Best Local Similarity
Matches 15; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; ORGANISM: Bovine US-60-252-833-24181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: RNA
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09/157,100

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APPLICANT: Sidransky, David
TITLE OF INVENTION: DETECTION OF NEOPLASIM BY ANALYSIS OF SALIVA
NUMBER OF SEQUENCES: 195
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 4225 Executive Square, Suite 1400
CITY: La Jolla
STATE: CA
COMPUTRY: USA
ZIP: 92037
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                            COMPUTER REALMALL

MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OCMPUTER: TBM Compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: FASTSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/863,806
FILING DATE: 12-May-1201
APPLICATION NUMBER: 09/038,637
FILING DATE: 12-NOV-1993
ATTORNEY AGENT INFORMATION:
NAME: Haile, Lisa A:
REGISTRATION NUMBER: 08/152,313
ATTORNEY AGENT INFORMATION:
NAME: Haile, Lisa A:
REGISTRATION NUMBER: 38,347
RELECOMMUNICATION INFORMATION:
TELEPHONE: 619/678-5570
TELEPHONE: 619/678-5570
TELEPHONE: 619/678-5570
TELEPHONE: 619/678-5570
TELEPHONE: 619/678-5099
INFORMATION FOR SEQ ID NO: 69:
SEQUENCE CHARACTERISTICS:
LENGTH: 26 base pairs
TYPE: INCIPIC acid
STRANDEDNESS: SINGLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TOPOLOGY: linear
; MOLECULE TYPE: Genomic DNA;
; SEQUENCE DESCRIPTION: SEQ ID NO: 69:
US-09-863-806-69
Sequence 69, Application US/09863806
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Db 23 CAAGTCAGTCTTA 11
Search completed: October 2, 2001, 16:55:02
Job time: 17665 sec

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0; Gaps

57.0%; Score 11.4; DB 6; Length 26; 92.3%; Pred. No. 1.3e+04; tive 0; Mismatches 1; Indels

Query Match 57.0 Best Local Similarity 92.3 Matches 12; Conservative

7 caggtgagtctta 19 || || || || || || || || || ||

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